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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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US-08-116-247-5
US-09-795-515-5
US-09-348-224-5
US-09-518-869-8
US-09-518-869-8
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US-09-526-738A-2
US-09-171-945-9
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US-09-171-769A-63
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29, Appli

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24, Appli

34, Appli

33, Appli

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36, Appli

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     Sequence
US-09-423-439-51

US-09-948-004-28

US-09-948-004-18

US-09-318-786-29

US-08-116-778E-2

US-08-438-562-2

US-08-438-562-2

US-08-438-562-2

US-08-438-562-2

US-08-438-561-3

US-08-434-122-3

US-09-434-122-33

US-08-308-494A-19

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Therapeutic Use in
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USING SINGLE CHAIN
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19;
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Pred. No. 1.2e-41;
Mismatches 19
RESULT 1

US-09-171-945-17

Sequence 17, Application US/09171945

Patent No. 6277599

GENERAL INFORMATION:
APPLICANT: Emery, Stephen
APPLICANT: Edge, Michael Derek
TITLE OF INVENTION: Monoclonal Antibody to CEA
TITLE OF INVENTION: Antibody, and Their Therap
FILE REFERENCE: Monoclonal Antibody to CEA
CURRENT APPLICATION NUMBER: US/09/171,945

CURRENT APPLICATION NUMBER: GB9703103.3

PRIOR FILING DATE: 1997-02-14

PRIOR APPLICATION NUMBER: GB9609405.7

PRIOR FILING DATE: 1997-04-29

NUMBER OF SEQ ID NOS: 131

SEQ ID NOS: 131

SEQ ID NO 17

FEMALE OF SEQ ID NOS: 2.1
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5455030-3
;Patent No. 5455030
; APPLICANT: LADNER, ROBERT C.;BIRD, RC
; TITLE OF INVENTION: IMMUNOTHERAPHY US
;POLYPEPTIDE BINDING MOLECULES
; NUMBER OF SEQUENCES: 24
; CURRENT APPLICATION DATA:
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ilarity 76.9%;
Conservative
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TYPE: PRT
ORGANISM: Artificial Sequence
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90; Conserv
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ZIP: 19103

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/303,569B

TT.TNG DATE: 07-SEP-1994
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CITY: Philadelphia
STATE: PA
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Pred. No. 3.1e-40;
10; Mismatches 19
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Pred. No. 2.8e-40
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APPLICATION NUMBER: US/08/40,440
FILING DATE: 1-APR-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                 SSULT 3
5-08-303-569B-5
Sequence 5, Application US/08303569B
Patent No. 5859205
GENERAL INFORMATION:
APPLICANT: Adair, John R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
                                      APPLICATION NUMBER: 512,910 FILING DATE: 25-APR-1990 APPLICATION NUMBER: 299,617 FILING DATE: 19-JAN-1989 APPLICATION NUMBER: 92,110 FILING DATE: 02-SEP-1987 APPLICATION NUMBER: 902,971 FILING DATE: 01-SEP-1986
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TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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larity 77.6%;
Conservative
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                                                                                                                                                                                                  APPLICANT: Jolliffe, Linda K.
APPLICANT: Jolliffe, Linda K.
APPLICANT: Zivin, Robert A.
APPLICANT: Adair, John R.
APPLICANT: Athwal, Diljeet S.
TITLE OF INVENTION: CD3 Specific Recombinant Ant NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Wasburn Kurtz Mackiewicz & STREET: One Liberty Place - 46th Floor CITY: Philadelphia
STATE: PA
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Pred. No. 3.1e-40;
); Mismatches 19;
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Floor
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                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 19103

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/116,247
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION ATA:
APPLICATION NUMBER: US 07/743,377
FILING DATE: 10-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Paintin, Francis A.
REGISTRATION NUMBER: 19,386
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CARP-0011
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; Sequence 5, Application US/08116247
; Patent No. 5929212
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
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ilarity 75.0%;
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GY: linear
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PRIOK L. SEQ NUMBER OF SEQ SEQ ID NO 8
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US-09-553-498-8
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LENGTH: 255
TYPE: PRT
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; ORGANISM: M
US-09-348-224-5
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Matches 88
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Best Local S
Matches 87
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Patent No. 6750325

GENERAL INFORMATION:
APPLICANT: Jollieffe, Linda Kay
APPLICANT: Zivin, Robert Allan
APPLICANT: Zivin, Robert Allan
APPLICANT: Athwal, John Robert
APPLICANT: Athwal, Diljeet Singh
TITLE OF INVENTION: CD3 Specific Recominant Antibody
FILE REFERENCE: CARPO066
CURRENT FILING DATE: 1999-07-06
EARLIER FILING DATE: 1999-07-06
BARLIER FILING DATE: 1993-09-03
EARLIER FILING DATE: 1991-10-04
NUMBER OF SEQ ID NOS: 29
COFTWARE: Patentin Ver. 2.1
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                                                                         . Kurtz Mackiewicz
46th Floor
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larity 75.0%; Pred. No. 3.1e-40,
Conservative 10; Mismatches 15
                                                                                                                                                                                                                        Version
                              Humanised Antibodies
                                                                                                                                 ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,515
                                                                                                                                                                                                                                                                                                                                                                                      CARP-0057
                                                                                                                                                                                                                                           .A:
US/09/795,515
                                                                                                                                                                                                                                                                     FILLING DATA:
CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/846,658
FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
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3-09-348-224-5
3-09-348-224-5
Sequence 5, Application US/09348224
Perent No. 6750325
                                                                      ADDRESSEE: Woodcock Washburn
STREET: One Liberty Place - 4
CITY: Philadelphia
STATE: PA
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TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
 APPLICANT: Athwal, Diljeet
APPLICANT: Emtage, John S.
TITLE OF INVENTION: Humanis
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
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nino acid
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Best Local Similarity
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Sequence 8, Application US/09553498

Sequence 8, Application US/09553498

Patent No. 6309861

GENERAL INFORMATION:
APPLICANT: Ambrosius, Dorothee
APPLICANT: Rudolph, Rainer
APPLICANT: Schaeffner, Joerg
APPLICANT: Schaeffner, Joerg
APPLICANT: Schwarz, Elisabeth
TITLE OF INVENTION: Process for the production of naturally folded and secreted
TITLE OF INVENTION NUMBER: US/09/553,498

CURRENT APPLICATION NUMBER: EP99107412.1
PRIOR APPLICATION NUMBER: EP99107412.1
PRIOR FILING DATE: 1999-04-26
NUMBER OF SEQ ID NOS: 10
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GENERAL INFORMATION:

APPLICANT: Ambrosius, Dorthee

APPLICANT: Rudolph, Rainer

APPLICANT: Schwarz, Elisabeth

TITLE OF INVENTION: PROCESS FOR THE PRODUCTION OF MOLECULAR

TITLE OF INVENTION: CHAPERONES

TITLE OF INVENTION: CHAPERONES

FILE REFERENCE: 20381

CURRENT APPLICATION NUMBER: US/09/618,869

CURRENT PILING DATE: 1999-07-19

PRIOR FILING DATE: 1999-07-29

NUMBER OF SEQ ID NOS: 10

SOFTWARE: PALENTIN Ver. 2.1
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                                                                     IELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPARL
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Length 235;
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4.4e-40;
  ore 445; DB 4; ]
ed. No. 3.1e-40;
Mismatches 19;
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Pred.
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illarity 75.0%;
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SOFTWARE: Pactor of the Software of the
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US-09-526-738A-2
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US-09-171-945-9
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Patent No. 5885816
GENERAL INFORMATION:
APPLICANT: Ikuo FUJII et al.
TITLE OF INVENTION: CATALYTIC ANTIBODIES ENANTIOSELECTIVELY
TITLE OF INVENTION: HYDROLYSING AMINO ACID ESTER DERIVATIVES
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, Lind & Ponack
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                                                                                                                     Indels
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                                                                        Length
                                                                                                                   15;
                                                                  core 444; DB 4;
ced. No. 4.4e-40;
Mismatches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73.2%; Score 443.5; DB 2;
larity 73.5%; Pred. No. 4e-40;
Conservative 12; Mismatches 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Wenderoth, Lind & Ponack STREET: 805 Fifteenth Street, N.W., #7 CITY: Washington STATE: D.C. COUNTRY: U.S.»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOMBER: US/08/737,129A
No. 5885816ember 15, 19
No. 435
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                                                                     Score
Pred.
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FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TTELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.5 in COMPUTER: IBM Compatible OPERATING SYSTEM: MS-DOS SOFTWARE: Wordperfect 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/73
                                                                                                               .;
                                                             73.3%;
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; ORGANISM: Escherichia coli
US-09-618-869-8
                                                                                                                Conservative
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SEQUENCE CHARACTERISTICS
LENGTH: 213 amino aci
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APPLICATION NUMBER:
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U.S.A.
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88; Conser
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                                                                    Query Match
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Sequence 2, Application US/09526738A
Sequence 2, Application US/09526738A
Patent No. 6630584
GENERAL INFORMATION:
APPLICANT: RAMOT UNIVERSITY AUTHORITY FOR APPLIED RESEARCH & INDUSTRIAL DVELOPMENT
APPLICANT: LTD.
TITLE OF INVENTION: SINGLE CHAIN ANTIBODY AGAINST MUTANT P53
FILE REFERENCE: 1196336
CURRENT APPLICATION NUMBER: US/09/526,738A
CURRENT FILING DATE: 2000-03-16
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
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                                                                                                       Comprising Said
an Adept System
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                                           APPLICANT: Emery, Stephen
APPLICANT: Copley, Clive Graham
APPLICANT: Edge, Michael Derek
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjuge
TITLE OF INVENTION: Antibody, and Their Therapeutic Use
FILE REFERENCE: Monoclonal Antibody to CEA
CURRENT APPLICATION NUMBER: US/09/171,945
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: GB9703103.3
PRIOR FILING DATE: 1996-05-04
PRIOR FILING DATE: 1996-05-04
PRIOR FILING DATE: 1997-04-29
MIMBER OF SEA TO NOTE
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 439; DB 3;
Pred. No. 5.1e-40;
; Mismatches 16
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Pred. No. 3e-38;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                          of Artificial
Sequence 9, Application US/09171945
Patent No. 6277599
GENERAL INFORMATION:
APPLICANT: Emery, Stephen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
9
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Similarity 79.6%;
86; Conservative 6
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4; Conservative
                                                                                                                                                                                                                                                                                        131
                                                                                                                                                                                                                                                                                                    Patentin Ver.
                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin V
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LTOSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPARLSG
                                                                         SGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRA 108
                                                                                                                                                                                                                                    Sequence;
Patent No. 5837245

Patent No. 5837245

GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Chezian Somasundaram
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, Suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRA
                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,052
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5
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426; DB 2;
No. 4.2e-38
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                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/484,172
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-043CP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
  Score Pred. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM PC compatible ISM PC compatible ISMS-DOS
                                                                                                                                                                                                                                   Sequence 16, Application US/0866105. Patent No. 5837243
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70.3%;
ilarity 79.0%;
Conservative
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SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acids
TYPE: amino acid
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC COM
OPERATING SYSTEM: PC
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83; Conserv
                   Similarity
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                                                                                                                                                                                                          RESULT 14
US-08-661-052-16
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                                 83;
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Best Local S
Matches 83
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                                            Sequence 4, Application US/09526738A
; Sequence 4, Application US/09526738A
; Patent No. 6630584
; GENERAL INFORMATION:
; APPLICANT: LTD.
; TITLE OF INVENTION: SINGLE CHAIN ANTIBODY AGAINST MUTANT P53
; FILE REFERENCE: 1196336
; CURRENT APPLICATION NUMBER: US/09/526,738A
; CURRENT FILING DATE: 2000-03-16
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                       Length 258;
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Pred. No. 3.1e-38;
; Mismatches 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nixon & Vanderhye, P.C.
30 No. 5876691th Glebe Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,507
FILING DATE: 02-Jul-1996
CLASSIFICATION: 530
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SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08652507
Patent No. 5876691
GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Arthur R. Crawford
REGISTRATION NUMBER: 25,327
REFERENCE/DOCKET NUMBER: 117
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4000
TELEFAX: 703-816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25,327
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iilarity 73.7%;
Conservative
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ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: AN' TITLE OF INVENTION: NUMBER OF SEQUENCES: 8 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  270 amino
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC (OPERATING SYSTEM: SOFTWARE: Patenti
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: USA
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84; Conserv
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US-08-652-507-2
                                                                                                                                                                                                                                                                             ORGANISM: Humanus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                             -09-526-738A-4
                                                                                                                                                                                                                                                258
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Best Local
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Matches
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US-09-188-082-16
Sequence 16, Application US/09188082
GENERAL INFORMATION:
APPLICANT: Yashwant M. Deo
APPLICANT: Yoshwant M. Deo
APPLICANT: Chezian Somasundaram
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISES
INVESTED OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSE: LAHIVE & COCKFIELD
STREET: Boston
STATE: Massachusetts
COUNTRY: USA
ZIP: O2109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BARDABLE FORM:
MEDIUM TYPE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/188,082
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/661,052
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAMME: ATTORNEY/AGENT INFORMATION:
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Pred. No. 1e-37;
7; Mismatches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MXI-043CP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: linear
NOLECULE TYPE: protein
US-09-188-082-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7;
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Best Local Similarity 79.0%;
Matches 83; Conservative
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Appliant App

Sequence Sequence

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Title: Perfect

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CELL POPULATIONS
US-09-189-129-3

US-09-824-286-3

US-08-860-174A-2

US-08-483-749A-14

US-08-621-751A-4

5455030-13

US-08-554-765-13

US-08-397-411-4

US-08-756-416-39

US-08-477-15

US-08-477-15

US-08-470-110A-15

US-08-881-037-66

US-08-891-196-4

US-08-752-844-66

US-09-293-533-4

US-09-293-533-6

US-09-293-533-6

US-08-652-558-8

US-08-652-558-8

US-08-122-546-12

US-09-131-052-12

US-09-131-052-12

US-09-131-053A-12

US-09-131-053A-12

US-09-131-053A-14

US-08-564-164A-4
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US-08-190-199A-67

i Sequence 67, Application US/08190199A

i Patent No. 5830663

j GENERAL INFORMATION:

APPLICANT: EMBLETON, Michael J.

APPLICANT: JONES, Peter T.

APPLICANT: WINTER, GREGORY P.

TITLE OF INVENTION: TREATMENT OF CELL

NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:

ADDRESSEE: PILLSBURY MADISON & SUTRC
STREET: 1100 New York Avenue, N.W.

CITY: Washington
STATE: D.C.

COUNTRY: U.S.A.

ZIP: 20005-3918

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/190,199A

FILING DATE: 13-JUL-1994

CLASSIFICATION NUMBER: PCT/GB92/01483

FILING DATE: 10-JUG-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: GB 9212419.7

FILING DATE: 11-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E SUTRO,
N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     льк: GB 9117352.6
10-AUG-1991
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     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 10-AUG-
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       Command line parameters:
-MODEL=frame+ n2p.model -DEV=xlp
-Q=/cgn2_1/USPTO_spool_p/US10049868/runat_13102004_132818_24818/app_query.fasta_1.1038
-Q=/cgn2_1/USPTO_spool_p/US10049868/runat_13102004_132818_24818/app_query.fasta_1.1038
-DB=Issued_Patents_AA -QFMT=fastan -SUFFIX=rai -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0_-UNITS=Dits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US10049868 @CGN_1_1 66 @runat_13102004_132818_24818 -NCPU=6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-DEV_TIMEOUT=120 -WARN_TIMEOUT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 67, Appl
Sequence 38, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 4, Appli
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Sequence 67, Appli
Sequence 67, Appl
Sequence 15, Appl
                                                                                                                                     updates/sec
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Compugen Ltd.
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US-08-190-199A-61
US-08-652-558-38
US-08-752-844-16
US-08-591-196-16
US-08-591-196-16
US-08-943-136-4
US-08-973-518-4
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US-08-881-037-67
US-08-667-769A-15
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Maximum Match 100%
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Indels:
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Batent No. 5830663

GENERAL INFORMATION:

APPLICANT: EMBLETON, Michael J.

APPLICANT: GOROCHOV, GUY

APPLICANT: JONES, Peter T.

APPLICANT: WINTER, Gregory P.

TITLE OF INVENTION: TREATMENT OF CEI

NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:

ADDRESSEE: PILLSBURY MADISON & SU

STREET: 1100 New York Avenue, N.W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/190,199A FILING DATE: 13-JUL-1994 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005-3918
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 2
US-08-190-199A-61
; Sequence 61, Application US/08190199A
; "---nt No. 5830663
  67
                                                                                1.62e-59
538.00
87.60%
85.95%
79.94%
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 222 amino acids
TYPE: amino acid
                                                protein
                                                                                                                                               (1-363)
                                     linear
                                                                                       Score:
Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                       113
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                                             ; MOLECULE TYPE:
US-08-190-199A-67
                                                                        Alignment Scores:
Pred. No.:
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                          TYPE: ami
TOPOLOGY:
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US-08-652-558-38
; Sequence 38, Application US/08652558
; Patent No. 5861155
; GENERAL INFORMATION:
; APPLICANT: LIN, AUGUSTINE YEE-THARN
; TITLE OF INVENTION: THEREOF
; NUMBER OF SEQUENCES: 52
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: BANNER & WITCOFF
STREET: 75 STATE STREET, 23RD FLOOR
                                                                                                                                                                                             Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette, 1.4 Mb
                                                                                                                                                                                                                                                                      x US-08-190-199A-61
                                                                                                                                                                                                                                                 Gaps:
PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/GB92/01483
FILING DATE: 10-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9212419.7
FILING DATE: 11-JUN-1992
PRIOR APPLICATION DATA:
                                                                            œ.
                                                                        APPLICATION NUMBER: GB 9117352
FILING DATE: 10-AUG-1991
INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
TYPE: amino acid
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538.00
87.60%
85.95%
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CITY: BOSTON
STATE: MASSAC
COUNTRY: USA
ZIP: 02109
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Pred. No.:
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3-08-752-844-16
Sequence 16, Application US/08752844
Patent No. 5935821
GENERAL INFORMATION:
APPLICANT: Chatterjee, Malaya
APPLICANT: Foon, Kenneth A.
APPLICANT: Chatterjee, Sunil K.
APPLICANT: Chatterjee, Sunil K.
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CELL CARCINOMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               120
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                  Length:
Matches:
                                     APPLICATION NUMBER: US/08/652,558
FILING DATE: JUNE 6, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/IB94/00387
FILING DATE: NOVEMBER 21, 1994
ATTORNEY/AGENT INFORMATION:
NAME: YANKWICH, LEON R.
REGISTRATION NUMBER: 30,237
REFERENCE/DOCKET NUMBER: 95,497-L
TELEPHONE: 617-345-9111
                                                                                                                                                                                                                                                                                                                                        Gaps:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                        38:
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535.50
90.91%
85.95%
79.57%
                                                                                                                                                                                             120 amino acids amino acids
                                                                                                                                                            TELEFAX: 617-345-9111
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 120 amino acid
TYPE: amino acid
                                                                                                                                                                                                               STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
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Conservative:
Mismatches:
Indels:
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NUMBER OF SEQUENCES: 66

CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
CITY: DALO ALTO
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/752,844
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30414-20002.21
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 494-0792
TELEFX: 706141
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                         59
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535.00
88.43%
85.95%
79.49%
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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
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Best Local Similarity:
Query Match:
DB:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 119 amino acid
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: sing
TOPOLOGY: linear
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Best Local Similarity:
Query Match:
                                                   363
                                                                           119
                                                                                                                                                                                                                                                                                           USA
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US-09-293-533-16
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US-09-293-533-16
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CITY: PA
STATE: C
COUNTRY:
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S-08-591-196-16
Sequence 16, Application US/08591196
Patent No. 5977316
GENERAL INFORMATION:
APPLICANT: Chatterjee, Malaya
APPLICANT: Foon, Kenneth A.
APPLICANT: Chatterjee, Sunil K.
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR 7
TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CELL
NUMBER OF SEQUENCES: 57
                                                                                                                                                                           COUNTRY: USA
ZIR: 94304-1018
ZIR: 94304-1018

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/591,196
FILING DATE: 16-JAN-1996
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30414-20002.20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1119
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Conservative:
Mismatches:
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Matches:
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ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
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535.00
88.43%
85.95%
79.49%
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity
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US-08-591-196-16
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FOR THE CELL CARCINOMA
                                                                                                                                                            Sequence 16, Application US/09293533
; Sequence 16, Application US/09293533
; Patent No. 6509016
; GENERAL INFORMATION:
; APPLICANT: Chatterjee, Malaya
; APPLICANT: Chatterjee, Sunil K.
; TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR
; TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CEL
; NUMBER OF SEQUENCES: 66
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 PAGE MILL ROAD
; CITY: PALO ALTO
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Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/293,533
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/752,844
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30414-20002.21
TELEPHONE: (415) 813-5600
TELEPHONE: (415) 494-0792
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Matches:
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                                         x US-08-943-136-4 (1-140)
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6th Floor, 330 University Avenue
Indels:
              Gaps:
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CITY: 10.

STATE: Ontario

COUNTRY: Canada

ZIP: M5G 1R7

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: OPERATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,518
FILING DATE: 07-APR-1998
CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/08973518
; Sequence 4, Application US/08973518
; GENERAL INFORMATION:
; APPLICANT: Anand, Naveen N
APPLICANT: Barber, Brian H
APPLICANT: Cates, George A
APPLICANT: Caterini, Judith E
APPLICANT: Caterini, Judith E
APPLICANT: Klein, Michel H
TITLE OF INVENTION: ANTIGENS TO SI
; TITLE OF INVENTION: ANTIGENS TO SI
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Sim & McBurney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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amino acid
78.97%
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                                           (1-363)
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Sequence 4, Application US/08943136

Patent No. 6291208

GENERAL INFORMATION:

APPLICANT: Anand, Naveen N

APPLICANT: Cates, George A

APPLICANT: Caterini, Judith E

APPLICANT: Klein, Michel H

TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR DELIVERY OF

TITLE OF INVENTION: ANTIGENS TO SELECTED CELLS OF THE IMMUNE
NUMBER OF SEQUENCES: 20

CORRESPONDENCE ADDRESS:
                                                        140
103
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Mismatches:
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6th Floor, 330 University Avenue
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Matches:
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STREET: 6th Floor, 330 University P
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/943,136
FILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION: 536
FILING DATE:
APPLICATION NUMBER: US 08/483,576
FILING DATE: 07-JUNE-1995
CLASSIFICATION: 536
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NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 140 amino acids
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531.50
88.43%
85.12%
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Best Local Similarity
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Pred. No.
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REFERENCE/DOCKET NUMBER: 203442110710
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
 5455030-15
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SEQUENCE CHARACTERISTICS:
LENGTH: 121 amino acid
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 (1-363)
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 -10-049-868A-2
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US-08-881-037-6
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5455030-15
5 Patent No. 5455030
7 TITLE OF INVENTION: IMMUNOTHERAPHY USING SINGLE CHAIN
7 POLYPEPTIDE BINDING MOLECULES
7 NUMBER OF SEQUENCES: 24
7 CURRENT APPLICATION DATA:
7 APPLICATION NUMBER: US/08/40,440
7 FILING DATE: 1-APR-1993
7 FILING DATE: 25-APR-1990
7 FILING DATE: 25-APR-1990
7 FILING DATE: 19-JAN-1989
7 FILING DATE: 19-JAN-1989
7 FILING DATE: 02-SEP-1987
7 FILING DATE: 02-SEP-1987
7 FILING DATE: 01-SEP-1986
7 FILING DATE: 01-SEP-1986
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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Matches:
Conservative:
Mismatches:
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85.12*
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529.50
90.00%
85.83%
78.68%
singl
                                                                                                -363)
STRANDEDNESS: Sil
TOPOLOGY: linear
-08-973-518-4
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Percent Similarity:
Best Local Similarity
                                                      Percent Similarity:
Best Local Similarity
Query Match:
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5455030-15
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Pred. No.:
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US-08-881-037-67

Sequence 67, Application US/08881037

Patent No. 6080588

GENERAL INFORMATION:

APPLICANT: Glick, Gary D.

APPLICANT: Swanson, Patrick C.

TITLE OF INVENTION: DNA BINDING ANTIBODIES

NUMBER OF SEQUENCES: 113

CORRESPONDENCE ADDESS:

ADDRESSEE: Morrison & Foerster

STREET: 755 Page Mill Road

CITY: Palo Alto

STATE: CA

COUNTRY: USA

ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TRYE: FIOPPY disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US 08/443,540

FILING DATE: 13-WAY-1995

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: KONSK!, Antoinette P:

REFERENCE/DOCKET NUMBER: 203442110710
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Batent No. 5783184

Batent No. 5783184

GENERAL INFORMATION:

APPLICANT: Appelbaum, Edward R.

APPLICANT: Cook, Richard M.

APPLICANT: Gross, Mitchell S.

APPLICANT: Holmes, Stephen D.

APPLICANT: Theisen, Timothy W.

ITILE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders NUMBER OF SEQUENCES: 76

CORRESPONDENCE ADDRESS:

ADDRESSEE: SmithKline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220

CITY: King of Prussia STATE: Pennsylvania
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114
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Matches:
Conservative:
Mismatches:
Indels:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/667,769A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17082
FILING DATE: 22-DEC-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/470110
525.50
87.60%
85.95%
78.08%
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CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d
                                                 US-10-049-868A-2 (1-363)
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CLASSIFICATION:
      Percent Similarity:
Best Local Similarity
Query Match:
DB:
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Matches:
Conservative:
Mismatches:
Indels:
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PCT-US95-17082A-15
; Sequence 15, Application PC/TUS9517082A
; GENERAL INFORMATION:
; APPLICANT: Ames, Robert S.
; APPLICANT: Appelbaum, Edward R.
; APPLICANT: Chaiken, Irwin M.
; APPLICANT: Cook, Richard M.
; APPLICANT: Gross, Mitchell S.
; APPLICANT: Holmes, Stephen D.
; APPLICANT: Holmes, Stephen D.
; APPLICANT: Theisen, Timothy W.
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50503
TELEPHONE: 610-270-5024
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
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Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
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523.00
88.43%
82.64%
77.71%
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Percent Similarity:
Best Local Similarity
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Pred. No.:
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DB:
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STRANDEDNESS:
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FRAGMENT TYPE:
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Mismatches:
Indels:
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                                                                                                                                                         Version
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Matches:
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                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US OR/10110

FILING DATE:
                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ve

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US95/170821
                                                                                                                                                                                                                                                                                                                       NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                          us 08/363131
Recombinant II
Treatment of :
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                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 0:
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
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523.00
88.43%
82.64%
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                                            SEE: SmithKline F: P.O. Box 1539-t
King of Prussia
                                                                                                                                                                                                                                                                                                                                                                             XX: 610-270-5090
ON FOR SEQ ID NO:
TITLE OF INVENTION: ReccTITLE OF INVENTION: TresNUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline
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Best Local Similarity
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PCT-US95-17082A-15
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TOPOLOGY: li
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CITY: 1
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US-09-189-3

US-09-189-129-3

Sequence 3, Application US/09189129

Patent No. 6323027

GENERAL INFORMATION:

APPLICANT: Burkly, Linda C

APPLICANT: Hession, Catherine A

APPLICANT: Whitty, Adrian

TITLE OF INVENTION: COMMON GAMMA CHAIN BLOCKING AGENTS

NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:

ADDRESSEE: Biogen, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1112
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Version
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/189,129
FILING DATE:
APPLICATION NUMBER: A006 PCT CIP
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 60/017,466
FILING DATE: 10-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan, Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A006 PCT C
TELEPHONE: 617 679-2000
TELEFAX: 617 679-2838
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCT
                                                                                                                                                                                                                                                                       STATE: Massachusetts
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : 112 amino acids
amino acid
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521.50
92.17%
85.22%
77.49%
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internal
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Best Local Similarity
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Matches:
Conservative:
Mismatches:
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ANALOGUE
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N.W.
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; Sequence 2, Application US/08860174A
; Patent No. 5989830
; GENERAL INFORMATION:
; APPLICANT: DAVIS, Paul James
APPLICANT: VERHOEIJEN, Martine Elisa
; APPLICANT: WILSON, Steve
; TITLE OF INVENTION: A BIFUNCTIONAL OR BITTILE OF INVENTION: ANTIBODY FRAGMENT AND NUMBER OF SEQUENCES: 31
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: PILLSBURY, MADISON & SUTRO STREET: 9th Floor, East Tower
CITY: WASHINGTON, D.C.
; STATE:
; COUNTRY: UNITED STATES
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FILING DATE: October 16, 1995
APPLICATION NUMBER: PCT/EP/96/03605
FILING DATE: August 14, 1996
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                       US-09-824-286-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS DC
SOFTWARE: MS WORD, WINDOWS 95, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,174A
FILING DATE: June 16, 1997
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YSTEM: PC-DOS/MS-DOS
MS WORD, WINDOWS 95, 1
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ZIP: 20005-3918
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy dis
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/824,286
FILING DATE: 02-Apr-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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NAME: Kaplan, Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A006 PCT CIP
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FILING DATE: 10-MAY-1996
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Benjamin, Christopher D
Hession, Catherine A
Whitty, Adrian
TITLE OF INVENTION: COMMON GAMMA CH
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
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STREET: 14 Cambridge Center
CITY: Cambridge
STATE: Massachusetts
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TELEPHONE: 617 679-2000
TELEFAX: 617 679-2838
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MEDIUM TYPE: Floppy disk
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TOPOLOGY: linear
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FRAGMENT TYPE: internal
SEQUENCE DESCRIPTION: SEQ
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SEQUENCE CHARACTERISTICS
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Query Match: DB:	ch:	76.89% 2	Indels: Gaps:	5	
US-10-049	0-049-868A-2 (1-363)	3) \times US-08-860-174A-2	74A-2 (1-239)		
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qq	124 GlnValGl	nLeuGlnGluSerGly	ProglyLeuValAla	Proserginserbenserile 143	
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qa	144 ThrCysThrVals		iii LeuThrGlyTyrGly		
δλ	121 CCAGGAAA	GGGTCTGGAGTGGCTG	GGAGTAATATGGACT	GGTGGAAGCACAAATTATAAT 180	
qq	164 ProGlyLy		:::: GlyMetIleTrpGly	}	
δλ	181 TCGGCTCT	CATGTCCAGACTGAGC	ATCAGCAAAGACAAC	TCGGCTCTCATGTCCAGACTGAGCATCAGCAAGACAACTCCAAGAGCGAAGTTTTCTTA 240	
qa	184 SerAlaLe	 			
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qa	204 LysMetAs				
QY	301 ACTATGAT	TACGGCCTATGCTATG	GACTACTGGGGCCAA	GGGACCACGGTCACCGTCTCC 360	
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S25058
Ig kappa chain - mouse
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #
C;Accession: S25058
R;Fischer, R.; Voss, A.; Niersbach, M.; Munziker, W
submitted to the EMBL Data Library, July 1992
A;Description: Production of a Tobacco mosaic virus
A;Reference number: S25057
A;Accession: S25058
A;Accession: S25058
A;Keference: mRNA
A;Residues: 1-235 <FIS>
A;Cross-references: EMBL:X67211; NID:g54828; PIDN:C2
C;Superfamily: immunoglobulin V region; immunoglobu:F;38-111/Domain: immunoglobulin homology <IMM>
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A; Cross-references: GB:M27792; NID:g197159
A; Experimental source: mRNA clones for anti-phenyloxazolone antibody 6F6
A; Note: the authors translated the codon TTC for residue 8 as Pro and TTC for residue A; Note: the codon given for 1-Met (AGT) is inconsistent with the authors' translation R; Milstein, C.; Even, J.; Jarvis, J.M.; Gonzalez-Fernandez, A.; Gherardi, E.
Eur. J. Immunol. 22, 1627-1634, 1992
A; Title: Non-random features of the repertoire expressed by the members of one V kapp? A; Reference number: A49044; MUID:92289826; PMID:1601044
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779; A49044; B49044
Rocca-Serra, J.; Maekelae, O.
859-865, 1988
corial association of V genes: one VH gene co
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- mouse (fragment)
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                                                    in, H.; Su, C.Z.; Chen, C.J. J. 12, 648-653, 1996
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197 #sequence_revision
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19 kappa chain V regions (27.7.2 and 27.4b.2) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change 21-
C;Accession: A30562
R;Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattacharya
J. Immunol. 142, 888-893, 1989
A;Title: Amino acid substitutions in V-H CDR2 change the idiotype b
A;Reference number: A30562; MUID:89110066; PMID:2464031
A;Reference number: A30562, MUID:89110066; PMID:2464031
A;Accession: A30562
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-107 <SIK>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
       A; Experimental source: spleen and myeloma cell line MOPC 315.43
A; Note: sequence inconsistent with nucleotide translation
A; Note: sequence extracted from NCBI backbone (NCBIN:144174, NC
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: immunoglobulin
F; 16-89/Domain: immunoglobulin homology < IMM>
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A; Accession: A49044
A; Molecule type: DNA
A; Residues: 1-25 <MIL>
A; Residues: 1-25 <MIL>
A; Cross-references: GB: 837663; NID: 9250214; PIDN: AAB22331.1; PID: 925
A; Arcession: B49044
A; Molecule type: DNA
A; Residues: 114-116 <MIL2>
A; Accession: B49044
A; Molecule type: DNA
A; Residues: 114-116 <MIL2>
A; Cross-references: GB: 837664; NID: 9250215; PIDN: AAB22332.1; PID: 925
A; Experimental source: BALB/c germ-line
A; Mote: sequence extracted from NCBI backbone; partial nucleotide s
A; Mote: sequence extracted from NCBI backbone; NCBIN: 106807, NCBIP: 1 C; Genetics:
A; Mote: sequence extracted from NCBI backbone; nucleotide s
A; Note: sequence extracted from NCBI backbone; nucleotide s
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A;Molecule type: mRNA
A;Residues: 1-106 <AGA>
A;Cross-references: GB:S68985; NID:g545746; PIDN:AAB30096.1; PID
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C; Species: Mus musculus (house mouse)
C; Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change
C; Accession: B54378
R; Agazie, Y.M.; Lee, J.S.; Burkholder, G.D.
J. Biol. Chem. 269, 7019-7023, 1994
A; Title: Characterization of a new monoclonal antibody to triple
A; Reference number: A54378; MUID:94165109; PMID:7509814
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C; Species: Mus musculus (house mouse)
C; Date: 10-Jul-1998 #sequence_revision 10-Jul-1998 #text_change 21
C; Accession: PD0011
R; Deng, J.B.; Han, H.; Su, C.Z.; Chen, C.Q.
Chinese Blochem. J. 12, 648-653, 1996
A; Title: Generation of a phage display library of the immunoglobul A; Reference number: PC4405
A; Contents: Spleen
A; Molecule type: mRNA
A; Residues: 1-107 < DEN>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-89/Domain: immunoglobulin homology < IMM>
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Ig kappa chain V region - mouse
C; Species: Mus musculus (house mouse)
C; Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
C; Accession: $29591
R; Kavaler, J.
submitted to the EMBL Data Library, April 1991
A; Reference number: $26459
A; Accession: $29591
A; Reference number: $26459
A; Accession: $29591
A; Status: preliminary
A; Molecule type: mRNA
A; Status: preliminary
A; Molecule type: mRNA
C; Superferences: EMBL: X59094; NID: g52227; PIDN: CAA41820.1; PID: g52228
C; Superfemily: immunoglobulin V region; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
F; 15-88/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Date: 30-Sep-1991 #sequence revision 30-Sep-1991 #text_change 21-
C;Accession: S05268; JL0062; S03846
R;Levy, S.
submitted to the EMBL Data Library, February 1989
A;Reference number: S05267
A;Accession: S05268
A;Molecule type: mRNA
A;Residues: 1-132 < LEV>
A;Coss-references: EMBL:X14098; NID:g52562; PIDN:CAA32260.1; PID:g
A;Carroll, W.L.; Starnes, C.O.; Levy, R.; Levy, S.
J. Exp. Med. 168, 1607-1620, 1988
A;Reference number: JL0061; MUID:89035985; PMID:3141553
A;Reference number: JL0062, MUID:89035985; PMID:3141553
A;Residues: 1-120 < CAR>
A;Residues: 1-120 < CAR>
A;Cross-references: EMBL:X14098
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C; Species: Mus musculus (house mouse)
C; Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_chain C; Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_chain C; Date: 23-Mar-1989 #text_chain C; Accession: B30562
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhattid. J. Immunol. 142, 888-893, 1989
A; Title: Amino acid substitutions in V-H CDR2 change the idion A; Reference number: A30562; MUID:89110066; PMID:2464031
A; Reference number: A30562; MUID:89110066; PMID:2464031
A; Residues: preliminary
A; Molecule type: mRNA
A; Residues: 1-107 <SIK>
C; Superfamily: immunoglobulin V region; immunoglobulin homology cimmon F; 16-89/Domain: immunoglobulin homology <IMM>
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F;1-14/Domain: signal sequence (fragment) #status predicted <SIG>F;15-132/Product: Ig kappa chain (fragment) #status predicted <MAT>F;15-108/Domain: V region (V-kappa-1) <VRE>F;30-103/Domain: immunoglobulin homology <IMM>F;109-132/Domain: J region (J-kappa-4) (fragment) <JRE>
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C; Species: Mus musculus (house mouse)
C; Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_cha
C; Accession: PS0071
R; Levy, S.; Campbell, M.J.; Levy, R.
J. Exp. Med. 170, 1-13, 1989
A; Title: Functional immunoglobulin light chain genes are rep
A; Reference number: A92781; MUID:89310348; PMID:2501443
C; Superfamily: immunoglobulin V region; immunoglobulin homol
C; Keywords: heterotetramer; immunoglobulin
F; 16-89/Domain: immunoglobulin homology <IMM>
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S11119
Ig kappa chain V region (clone NQ6-8.3.1) - more; Species: Mus musculus (house mouse)
C; Species: Mus-1997 #sequence_revision 23-Aug-C; Accession: S11119
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F. Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually really Reference number: S07331; MUID:83271467; PMIA; Molecule type: mRNA
A; Residues: 1-107 <NAT>
C; Superfamily: immunoglobulin V region; immunogister immunoglobulin homology <IMM>
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Ig kappa chain V region (H18-S415) - mouse
C;Species: Mus musculus (house mouse)
C;Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 09
C;Accession: G27887
R;Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
EMBO J. 5, 1577-1587, 1986
A;Title: Structural and functional implications of a restricted an A;Reference number: A91043; MUID:86300658; PMID:2427335
A;Accession: G27887
A;Molecule type: DNA
A;Residues: 1-106 <CAT>
A;Residues: 1-106 <CAT>
A;Residues: 1-106 <CAT>
CCAT>
A;Residues: UNIPROT:Q8VDD0
A;Experimental source: strain Balb/c
A;Coss-references: UNIPROT:Q8VDD0
C;Comment: This chain was isolated from a hybridoma protein that b
C;Comment: This chain was isolated from a hybridoma protein that b
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 21
C;Accession: B49049
R;Armandola, E.A.; Mariani, S.M.; Zwickl, M.; Hardman, N.; Ferrone
Eur. J. Immunol. 22, 2893-2899, 1992
A;Title: Molecular analysis of anti-idiotypic monoclonal antibodie
A;Reference number: A49049; MUID:93049629; PMID:1425914
A;Accession: B49049
A;Accession: B49049
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
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Ig kappa chain V region (clone NQ5-61.1.2) - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_change 21-C; Accession: S11118
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG response A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Accession: S11118
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-107 < KAA>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 16-89/Domain: immunoglobulin homology < IMM>
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STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S., Ni
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ da
EMBL; AF516283; AAM64201.1; -.
PIR; H33932; H33932.
HSSP; Q91W12; 1AY1.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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KV6E MC
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Eukaryota; Metazoa; Chordata; Chammalia; Eutheria; Rodentia; SchorBI_TaxID=10090;
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-Q=/Cgn2_1/USPTO_spool_p/US10049868/runat_13102004_132817_24791/ap
-DB=UniProt_02 -QFMT=fastan -SUFFIX=rup -MINMATCH=0.1 -LOOPCL=0 -1
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -0
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-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=-
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STRUCE-Mammary tumor. WAP-TGF alpha model. 7 months old;

X STRAID=mix FVBAN;

X Strausberg R.L., Feingold E.A., Grouse b.H., Derge J.G.,

A Strausberg R.L., Seberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jozdan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Brownstein M.J., Uschiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S.A., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Nilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

A Nhiting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

A Rodriguez A.C., Grimwood J., Schwutz J., Myers R.M., Butterfield Y.S.,

A Jones S.J., Marra M.A.,

T "Generation and initial analysis of more than 15,000 full-length human
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                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mu
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences.";
id. Sci. U.S.A. 99:16899-16903(2002)
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                                                                                                                                                                                                                            ACCAAGCTGGAGCTGAAACGGGCTGATGCTGCACCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Mammary tumor, WAP-TGF alpha model. 7 Strausberg R.; Submitted (APR-2001) to the EMBL/GenBank/DDB.
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01-DEC-2001
01-MAR-2004
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Murinae; Mu
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Sciurognathi; Muridae;
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, Last sequence update)
, Last annotation update)
c chain (Fragment).
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Pfam; PF07654; C1-set; 1.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
Hypothetical protein.
SEQUENCE 235 AA; 26021 MW; 5FC73
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01-MAR-2002 (TrEMBLrel. 20, Cr
01-MAR-2004 (TrEMBLrel. 26, La
01-MAR-2004 (TrEMBLrel. 26, La
Anti-MOG Z12 variable light ch
Name=anti-MOG kappa;
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Submitted (OCT-2001) to the EM
EMBL; AJ416331; CAC94866.1; -.
PIR; G27887; G27887.
HSSP; P01834; 1MIM.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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STRAIN=BALB/c;
Sembi P.;
Submitted (JAN-2002)
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
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NCBI_TaxID=10090;

[1]

SEQUENCE FROM N.A.

STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;

Zhou Y.-X., Taguchi H., Planque S., Karle S., Nissubmitted (MAY-2002) to the EMBL/GenBank/DDBJ date EMBL; AF516285; AAM64203.1; -.

PIR; A33933; A33933.

PIR; PC4405; PC4405.

HSSP; P01837; 10RQ.

InterPro; IPR003596; Ig_v.

Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                  AGTGGGTCTGGGACAGAATTCACCCTGGAAATCAGTAGAGGTGAAGGCTGAGGATGTGGGT
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Sciurognathi; Muridae; Murinae; Mus
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Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity.
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MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein "mRNA sequences define an unusually restricted IgG rephenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody
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on update)
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P04940;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence upda
15-JUL-1999 (Rel. 38, Last annotation up
IS kappa chain V-VI region NQ2-17.4.1.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Eutheria; Rodentia; Sciurognat
NCBI_TaxID=10090;
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JASP; P016,

InterPro; IPR003596,

InterPro; IPR003596,

Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG LIKE; 1.

Hybridoma; Immunoglobulin V r

DOMAIN 24 33

DOMAIN 24 33

DOMAIN 34 48

SS 88 96
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Query Match:
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01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Musinae; MucBi TaxID=10090;
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databases.
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STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S., Nis
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ dat
EMBL; AF516284; AAM64202.1; -.
PIR; A33933; A33933.
PIR; PH1058; PH1058.
HSSP; P01837; 25C8.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                     Craniata; Vertebrata; Euteleostomi
Sciurognathi; Muridae; Murinae; Mu
                                                                                                                                                                                                                                                                                                                                                                                                                 PO4942;
PO4942;
PO4942;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
13-AUG-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation NQ5-61.1.2.

Mus musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; M
NCBI_TaxID=10090;

NCBI_TaxID=10090;

NCBI_TaxID=10090;

NCBI_TaxID=10090;

MEDLINE=83271467; PubMed=6877353;

Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;

MEDLINE=83271467; PubMed=6877353;

Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;

"mRNA sequences define an unusually restricted IgG response to phenyloxazolone and its early diversification.";

Nature 304:320-324(1983).

-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity.
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Hybridoma; Immunoglobulin V res
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[1]—SEQUENCE FROM N.A.
MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG response phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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                                                                                             GCTCGCCTCAGTGGCTCTGGGACAGAATTCACCCTGGAAATCAG
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                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Exthe European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-s or send an email to license@isb-sib.ch).
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Complementarity-determining-1
Framework-2.
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Mismatches:
Indels:
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13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain V-VI region NQ6-8.3.1.
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                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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HSSP; P01679; 2FBJ.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
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SEQUENCE FROM N.A.

STRAIN=CZECH II;

TISSUE=Mammary tumor metastatized to lung. MMTV-LTR/Wntl model.

TISSUE=Mammary tumor metastatized to lung. MMTV-LTR/Wntl model.

Expression driven by an MMTV-LTR enhancer.;

MEDLINE=22388257; PubMed=12477932;

A strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.W.,

B Brownstein M.J., Uschin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.W.,

A Nilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Boutferfield Y.S.

Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Jones S.J., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human "Generation and initial malysis of more than 15,000 full-length human "G.
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Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mu
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P04941;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IS kappa chain V-VI region NQ2-48.2.2.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus
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MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG response
                                                           l to lung. MMTV-LTR/Wntl
enhancer.;
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                                                                                           databases
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                                   SEQUENCE FROM N.A.
STRAIN=CZECH II;
TISSUE=Mammary tumor metastatized to lung. MMTV-I
Expression driven by an MMTV-LTR enhancer.;
Strausberg R.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ date EMBL; BC055911; AAH55911.1; -.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig-c1.
InterPro; IPR003596; Ig-c1.
InterPro; IPR003596; Ig-c1.
InterPro; IPR003596; Ig-c1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG-MHC; UNKNOWN_1.
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Matches:
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Eukaryota; Metazoa; C
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                            This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMI the European Bioinformatics Institute. There are no restriuse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-scor send an email to license@isb-sib.ch).
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Complementarity-determining-1
Framework-2.
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Complementarity-determining-3
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02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
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phenyloxazolone and its early diversifinature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone
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PROSITE; PS50835; IG LIKE;
Hybridoma; Immunoglobulin
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 Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Laxidae; Rodentia; Sciurognathi; Muridae; NCBI_TaxID=10090;
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Sciurognathi; Muridae;
                                    SEQUENCE FROM N.A.
STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhu "Antinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ da'
EMBL; AY436821; AAR10981.1; -.
NON TER 111 111
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STRAIN=B6.Slel; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K.,
"Antinuclear autoantibodies from B6.Slel mice.
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ
EMBL; AY436822; AAR10982.1; -.
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Sciurognathi; Muridae;
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AAR10994;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhuntinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ da
EMBL; AY436834; AAR10994.1; -.
Length:
Matches:
Conservative:
Mismatches:
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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AAR11048;
AAR11048;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi,
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; MucBI TaxID=10090;
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STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases
EMBL; AY436888; AAR11048.1; -.
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SEQUENCE 108 AA; 11738 MW; C11222F2FDFBC160 CRC64;
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New motif-grafted hybrid polypeptides binding to the infectious form of a prion, useful for diagnosing or treating diseases of protein aggregation or conformation, e.g. amyloidosis, Alzheimer's disease, renal failure or
    Ü
    Moroncini
     RA,
    Williamson
             WPI; 2003-877028/81.
N-PSDB; ADE06763.
                                          diabetes.
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English 34; 115pp; SEQ ID NO Disclosure;

The present invention describes a hybrid polypeptide (I) comprising: (a) a calf residues from a polypeptide associated with a disease of protein aggregation or conformation to bind an aggregating form of the polypeptide or to a disease-associate conformer of the polypeptide; and polypeptide or to a disease-associate conformer of the polypeptide; and polypeptide or to a disease-associate conformer of the polypeptide; or polypeptide binds with greater affinity to a disease causing or infectious conformer of the polypeptide that is the source of the polypeptide binds with greater affinity to a disease causing or infectious conformer of the polypeptide that is the source of the polypeptide motif compared to a benign form of the polypeptide. Also described: (I) a nucleic acid molecule; (3) a cell comprising the vector; (4) detecting an isoform or a Presc form of a prion polypeptide or a polypeptide associated with a disease of protein aggregation, (i) preparing a plyceptide or a polypeptide associated with a disease of protein aggregation, (7) preparing a hybrid molecule with a disease of protein aggregation, (7) preparing a hybrid molecule that specifically interacts with one conformer of a prion protein. (1) has neuroprotective, nootropic, antidiabetic, anti-idioxype antibody that cerebroprotective, antiparkinsonian, cytostatic, nephrotropic, antidiabetic, anticonvulsant, cerebroprotective, antiparkinsonian, cytostatic, nephrotropic, antidiabetic anti-idiabetic, and can be used in diagnosing or treating diseases of protein aggregation or conformation, such as Creutzfeld-Jakob disease, transinger amploidosis associated with chronic inflammatory gisease, receive amploidosis associated with chronic inflammatory disease, immunoglobulin amyloidosis, reactive amyloidosis associated with chronic inflammatory disease, immunoglobulin amyloidosis, reactive amyloidosis associated with chronic inflammatory disease, immunoglobulin amyloidosis, reactive amyloidosis associated with chronic inflammatory disease, immunoglobulin am copic, cardiant, I can be used in invention can be myeloma, plasma medullary carcinoma in a sample transthyretin gene, amyotrophic lateral sclerosis, Pick's disease, Parkinson's disease, Frontotemporal dementia, multiple myeloma, plasma cell dyscrasias, familial amyloidotic polyneuropathy, medullary carcinor of thyroid, chronic renal failure, congestive heart failure, senile cardiac and systemic amyloidosis, chronic inflammation, atherosclerosis or familial amyloidosis. The present sequence is used in the exemplification of the present invention.

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(1-215)x ADE06764 US-10-049-868A-1 (1-359)

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The present invention describes cell death-inducing agents containing the degradation product of an antibody which can recognise human leukocyte antigen (HLA). Also described: (1) a degradation product of the 2D7 antibody; (2) a similar degradation product of an antibody which is: (a) one containing an amino acid sequence of (SEQ ID NO:6) with 253 amino acids substituted, deleted, inserted and/or added and being functionally equivalent to the parent; (c) one containing the CDR amino-acid sequences of (SEQ ID NO:2) and (SEQ ID NO:2) and (SEQ ID NO:2) and (SEQ ID NO:4) but with some amino acids substituted, deleted, inserted and/or ID NO:4) but with some amino acids substituted, deleted, inserted and/or added and being functionally equivalent to the parent; (3) a process for producing an antibody with an increase of activity by degrading a HLA-recognising antibody, or by degradation products of antibody, inducing agents containing the degradation products of antibody, inducing acids inhibiting cell proliferation containing the degradation

Claim 6; SEQ ID NO 4; 106pp; Japanese.

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ThrSerProLysArgTrpIleTyrAspThrSerLysLeuAlaSerGlyValProAlaArg
                  CTCAGTGGCAGTGGGACTGGGAATTCACCCTGGAAATCAGTAGAGTGAAGGCTGAG
                                                               Cell death-inducing agents containing degradation product of antiboc
particularly diabody recognizing human leukocyte antigen (HLA),
applicable in antitumor agents and remedies for autoimmune diseases
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                                                                                          Mouse IgG2b light chain protein SEQ ID NO:4
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products of antibody, comprising the produced antibodies or 2D7 antibody, comprising the degradation products of antibody, comprising the produced antibodies or 2D7 antibody; and (7) remedies for autoimmune diseases containing the degradation products of antibody, comprising the produced antibodies or 2D7 antibody. The 2D7 antibody has cytostatic, immunosuppressive, antiinflammatory and dermatological activities. The cell death-inducing agents are applicable in antitumour agents and remedies for autoimmune diseases for treating e.g. cancer, Crohn's disease and systemic lupus erythematosus. The present sequence represents a mouse IgG2b light chain, which is used in the exemplification of the present invention.
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25-MAR
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                                                                                                                                            antibodies - used for as therapeutic agent
                                                                                                                                                                                                                                                                                                                                                       AGTGGCAGTGGGTCTGGGACAGAATTCACCCTGGAAATCAGTAGAGGTGAAGGCTGAGGAT
                                                                                                                                                                                          The MAD recognises the epitope on alpha/beta TCR and includes a med resonse. It is highly effective in clinical application before, aft during bone marrow transplant surgery, in treatment of cancer and autoimmune diseases, and has applications in immunoregulation. Abs also be used in detection of immunocompetent T-cells. (Updated on 2003 to correct PA field.)
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Matches:
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90US-00529979
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460.00
87.50%
80.36%
68.25%
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immuno-regulation, e.g. t
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BEHRINGWERKE AG
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N-PSDB; AAQ06956, A
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Best Local Similari
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25-MAY-1990;
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19H22 (isotype IgG2b/kappa). The mAb is produced by hybridoma cells obtained by immunising CB6 fl mice with human RANK-L protein, and fusing Spleen cells with X63-Ag8.653 myeloma cells. 19H22 mAb is specific for buman RANK-L, having a binding affinity characterised by a dissociation constant of about 10 power -10 M. RANK-L is a member of the tumour constant of about 10 power -10 M. RANK-L is a member of the immune system, bone development and homeostasis. The invention provides altered, chimeric and humanised 19H22 antibodies, including Fab or F(ab')2 fragments, antibodies comprising the light and heavy chain variable regions of 19H22. These are used in methods for the diagnosis, treatment and prevention of osteopenic diseases, including rheumatoid arthritis, osteoporosis, metastatic and primary bone cancer, wear debris induced osteologis, metastatic and primary bone cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis, treatment and prevention of osteopenic diseases, including rheumatoid arthritis, osteoporosis, metastatic and primary bone cancer, wear debris induced osteolysis or osteoarthritis, and immune diseases including psoriasis, insulin dependent diabetes, inflammatory bowel disease or multiple sclerosis. Methods are also provided for the recombinant production of the antibodies, using isolated nucleic acids and transfected host (preferably mammalian) host cells, and a method of diagnosing conditions associated with Th1 T-cell activity or osteoclast development and activation, in particular those listed above
                             complementarity determining region; CDR; rheumatoid arthritis; osteoporosis; bone cancer; metastasis; osteolysis; osteoarthritis; psoriasis; inflammatory bowel disease; multiple sclerosis; diabetes; cytostatic; antiarthritic; antiinflammatory; osteopathic; antidiabetic; antirheumatic; immunosuppressive; antipsoriatic; neuroprotective; light chain; VL; mouse.
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rheumatoid arthritis and osteporosis.
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               chain
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07-SEP-2000; 2000US-0230639P
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             Anti-RANK ligand mab
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N-PSDB; AE
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129

Length:

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Alignment Scores: Pred. No.:

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97GB-00003103
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The antibody is
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459.00
86.21%
76.72%
68.10%
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N-PSDB; AAV17288.
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Best Local Similarity:
Query Match:
DB:
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02-JUN-1998
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14-FEB-1997
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antibody (806.077 Ab). Host cells or transgenic organisms transformed with DNA encoding the antibody, are used to make the antibody or conjugate is used in a medicament suitable for intravenous administration. The conjugate can be used for cancer therapy, selectively killing tumour cells. The antibody can be used for in vivo or in vitro diagnosis of cancer. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                   240
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49. 55
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94. 96
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The VK sequence is one of 23 (AAR21286-308) expressed from a single chain fragments specific for 2-phenyl-5-oxazolone (phox). It was prepd. using cDNA generated from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by CCR (AAQ22474-84) and ligated into fdCAT2 (AAQ23463) for expression on the phage surface as fusions with gene III. The resulting library of clones was diverse; 23 hapten binding clones were sequenced revealing 8 vH genes (A-H; AAR21286-92). Most clones were VH-B combinations so a further hierarchical library was prepd. by "crossing" VH-B with the Vk repertoire. The resulting library was screened for hapten binding and 24 clones sequenced. 14 new partners (AAR21293-308) for VH-B were identified. Nearly all the Vk genes were "co.x-like"; only f, (from the identified Nearly all the Vk genes were "co.x-like"; only f, (from the original library) and h, p, q, and r (from the hierarchical library) were VkoxI type genes. Of the 24 hierarchical clones, 2 were of type "p". The Coriginal library and h, p, q, and r (from the hierarchical library) were the suggests that phage bearing scry fragments having weak affinit- ies can be selected with antigen, probably due to the avidity of the multiple antibody heads on the phage. The different combinations could also be isolated on a basis of antigen affinity. See also AAR21260-307, 309-311; AAR22450, 565-581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           120
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                                                                                                                               Hoogenboom HRJ,
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88
8
12
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (1-108)
                                                                                         ANTIBODY TECHNOLOGY
                                                                                                                              KS,
JD;
                                                                                                                                                                                                                                               English
                                                                                                                               Johnson
Marks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             x AAR21301
                                                                                                                                                                                               of specific cells with a
           90GB-00015198.
90GB-00022845.
90GB-00024503.
91GB-00004744.
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452.00
88.89%
81.48%
67.06%
                                                                                                      COUNCIL
                                                                                                                              J, Pope AR, Holliger
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                                                                                                                                                                    WPI; 1992-056862/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (1-359)
                                                                                         CAMBRIDGE
MED RES CO
                                                                                                                                                                                             Producing members recombinant host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Percent Similarity:
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 108 AA;
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           10-JUL-1990;
19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
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Pred. No.:
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Jackson RH,
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                                                                                         (CAMB-)
                                                                                                                                                                                                                       package
                                                                                                      (MEDI-)
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DB:
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GATGTGGGTGTGTATTACTGTCAACAACTTGTAGAGTATCCGCTCACGTTCGGTGCTGGG

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WPI; 1993-367875/46
N-PSDB; AAQ51535.
                                                                                                                                                                                                                                                            Mouse anti-bovine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     215
                                                                                                                                                                                                                                                                                                       Mus musculus
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02-SEP-1987;
19-JAN-1989;
                                                                                                                                                                                                                                  25-MAR-2003
23-MAY-1994
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                  21
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heavy c
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| AspvalAlaThrTyrTyrCysGlnGlnTrpSerSerAsnProLeuThrPheGlyAlaGly 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present
antibody BF11,
                                                                                                                                                                                                                                                                                                                                                                                                                         O
H
                                                                                                                                                                                                                                                                                                                                                                                                  The specification describes antibodies or polypeptides for microbial lipase, which enhance the hydrolytic activity of the enzyme. The polypeptides, antibodies or immunocomplex are useful for biocatalysis objectonversion reactions (e.g. hydrolysis, interesterification, esterification, alcoholysis, acidolysis or aminolysis); in biosensor technology; in bioremediation; in preparing fragrances or flavours and chemical modification of alcohols, acids, esters or fats; in cosmetic, pharmaceutical and food industries; in preparing detergents; or in identifying, purifying or measuring microbial lipases. The present sequence represents the variable light chain of monoclonal antibody BFI which is directed against the Candida rugosa lipase.
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                                                                                                                                                                                                                                                                                      Nuti
                                                                                                                                                                                                                                                                                                                                                            OL
                                                                                                                                                     flavour;
                                                                                                                                                                                                                                                                                                                                        New antibodies and polypeptides that recognize microbial lipases enhance the hydrolytic activity of the enzyme, useful for e.g. biocatalysis or bioconversion reactions, in biosensor technology
                                                                                                                                                                                                                                                                                      ß
                                                                                                                                    BF11
                                                                                                                                                                                                                                                                                      Barachini
                                                                                                                                                     biocatalysis; bioconversion; fragrance; chain; monoclonal antibody BF11.
                                                                                                                                    antibody
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18
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Mismatches:
Indels:
                                                                                                                                    chain
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                                                                                                                                    variable light
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                                                                                 standard; protein; 156
                                                                                                                                                                                                                                                                                      Rughetti
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                                                                                                                                                                                                                                                                                               Cernia E;
                                                                                                                                                                                                                                                                     STUDI
                                                                                                                                                                                                                                   2002WO-EP012149
                                                                                                                                                                                                                                                    31-OCT-2002; 2002WO-EP012149
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452.00
84.62%
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                                                                                                                                   acid sequence
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N-PSDB; ADO43519.
                                                                                                                   (first
                                                                                                                                                    antibody; lipase;
detergent; light
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Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 156 AA;
                                                                                                                                                                                                WO2004039845-A1
                                                                                                                                                                                                                                                                                                                                                                   bioremediation.
                                                                                                                                                                                                                                 31-OCT-2002;
                                                                                                                                                                               musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-049-868A-1
                                                                                                                   29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Scores
                                                                                                                                                                                                                 13-MAY-2004
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                                                                                                 ADO43520;
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           87
                            301
                                                                                 ADO43520
                                                                                                                                                                                                                                                                                               Palocci
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120
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                                                                                                                                                                                                                  80
                                          40
                                                                                                                              9
ATGACCTGCAGTGCCAGTTCAAGTGTTAATTACATGCACTGGTTCCAGCAGGAGTCGGGC
                                                                                  GATGTGGGTGTATTACTGTCAACAACTTGTAGAGTATCCGCTCACGTTCGGTGCTGGG
                                                                                                                                                                          CTCAGTGGCAGTGGGTCTGGGACAGAATTCACCCTGGAAATCAGTAGAGTGAAGGCTGAG
                                                                                                                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chain poly:peptide for binding antigen - comprising light chain antigen binding portions linked by peptide linker.
                                                                                                                                                                                                                                                                                                                                                 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosti
                                                                                                                                                                                                                                                                                                                                              antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               growth hormone MAb light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l antibody; MAb; affinity; binding; imaging; purification; biosensors.
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87US-00092110.
89US-00299617.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immuno:purificn, using single binding chain molecule includi
binding parts of antibody light and heavy chain variable reg
connected by a linker - is smaller, stabler and less expensi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antibody engineering; single polypeptide chain binding mheavy chain; light chain; monoclonal antibody; MAb; bovine growth hormone; bGH; immunoaffinity purification.
                                                            ive:
                  Length:
Matches:
Conservati
Mismatches
Indels:
Gaps:
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87US-00092110.
89US-00299617.
90US-00512910.
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(first en
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N-PSDB; AAT13734.
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                                                        ent Similarity:
Local Similarity
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02-SEP-1987;
19-JAN-1989;
25-APR-1990;
01-APR-1993;
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   Scores
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Alignment
Pred. No.:
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The mature heavy chain (AAR99643) and mature light chain (AAR99644) of the mouse anti-bovine growth hormone monoclonal antibody 3C2 can be utilised in novel single chain binding molecules (AAR99645-48), in which the hypervariable regions from IgG1 3C2 MAb are joined by peptide linkers derived from the Fv regions of an IgA class anti- phosphorylcholine myeloma antibody, MCPC-603. The single chain molecules retain the binding specificity of the light and heavy chains and have the advantages of smaller size, greater stability and reduced cost. They can be used in therapy, diagnostics, imaging, purification and biosensors. (Updated on 25-MAR-2003 to correct PF field.)
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 Portions of the heavy chain (AAR97376) and light chain (AAR97377) of murine IgG anti-bovine growth hormone monoclonal antibody 3C2 can be incorporated into novel single polypeptide chain binding molecules (see also AAW02188-90). These are expressed in host cells using DNA constructs (see also AAT36460-62) that include heavy and light chain encoding sequences (AAT29056 and AAT29057) joined by linker moieties. Following expression and refolding, the single chain binding molecules show the binding characteristics of the aggregate of the 2 original heavy and light chains of the variable region of the antibody. (Updated on 25-MAR-2003 to correct PF field.)
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The VK sequence is one of 23 (AAR21286-308) expressed from a single chain FY library. The library produces a diverse repertoire of antibody fragments specific for 2-phenyl-5-oxazolone (phox). It was prepd. using cDNA generated from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by CRR (AAQ23474-84) and ligated into fdCAT2 (AAQ2343) for expression on the phage surface as fusions with gene III. The resulting library of clones was diverse; 23 hapten binding clones were sequenced revealing 8 VH genes (A-H; AAR21264-71) in a variety of pairings with 7 VK genes (a-g; AAR21286-92). Most clones were VH-B combinations so a further hierarchical library was prepd, by "crossing" VH-B with the VK clones sequenced. 14 new partners (AAR21293-308) for VH-B were clones sequenced. 14 new partners (AAR21293-308) for VH-B were coriginal library) and h, p, q, and r (from the hierarchical library) were coriginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were coriginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library) and h, p, q, and r (from the hierarchical library) were voiginal library hierarchical librar
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g3p; binding; adsorption; gene VIII; diverse repertoire;
specific binding pairs; replicable genetic display package.
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90GB-00024503.
91GB-00004744.
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   Murine VL kappa
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Binding-site
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19-OCT-1990;
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recombinant
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                                                                                          The OK3T light chain sequence was deduced from the cDNA sequ
                                      New humanised antibodies comprising CDR grafted antibody light chains, for use in vivo therapy and diagnosis.
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Mismatches:
Indels:
Gaps:
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/label= signal peptide
23. .235
/label= light chain
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                          US-10-049-868A-1 (1-359) x AAR21293
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90WO-GB002017
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80.56%
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P-PSDB; AAR13060.
Similarity:
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21-DEC-1990;
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Query Match:
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from a library prepared from OK3T producing cells. The library was screened with a probe complementary to a region in the mouse kappa constant region. The OK3T sequence was used in CDR- grafting experiments to prepare humanised antibodies. NOTE: Data found in patent WO9109968 has been used to index this entry. (Updated on 25-MAR-2003 to correct PA field.)
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Matches:
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N-PSDB; ACC84876.
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The invention relates to increasing the content of one or more transgene-
coded biomolecules in an organism and involves changing the distribution
of ATP and/or ADP in cells of the organism. The yield of transgenic
molecules in host cells is often insufficient for industrial production.
The method increases the yield of transgenic molecules in animal and
plant host cells, therefore facilitating their production on an
industrial scale. The proteins produced by the method are useful for
diagnosing, preventing and/or treating viral diseases and cancer. The
present sequence represents a single-chain (scFv) antibody, used to
exemplify the increase in the expression of scFv antibodies in transgenic
potato tubers
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                  enting and/or distribution
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                                                                                                                                                                                                                                                                                                                                                                                  antibody; hormone;
increasing the content of transgene-coded biomolecules in animal, useful for producing proteins for diagnosing, prevetreating viral diseases and cancer, comprises changing the of ATP and/or ADP.
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                                                                                                                                                                                This invention describes a novel preparation of a water-soluble, naturally occurring eukaryotic polypeptide containing two or more cysteine units bound via a disulfide bridge which comprises cultivation of prokaryotic cells in the presence of arginine or an amide compound. The method is useful for the preparation of eukaryotic proteins e.g. proteases, interferons, protein hormones, antibodies or antibody fragments (e.g. a single chain FV fragment that binds to thyroid stimulating hormone). It is especially useful for preparing proteins with more than five disulfide bridges, e.g. recombinant plasminogen activator (rPA). The technique is simple and does not require in vitro after-treatment, such as the removal of inclusion bodies, reduction or naturization
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                                                                                                                   Preparation of water-soluble e
bridges e.g. rPA, comprises cu
presence of arginine or amide
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444.00
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N-PSDB; AAC66074.
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Best Local Similarity
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    PelB-scFvOxazolon fusion protein.
                                                   24-JUL-2000; 2000EP-00115839.
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             Molecular chaperone; PelB
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                                                                                Rudolph
                                                                                        WPI; 2001-246712/26.
N-PSDB; AAF77806.
                       Unidentified
                                 EP1077262-A1
                                                            29-JUL-1999;
                                                                               Ambrosius D,
                                          21-FEB-2001.
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The present invention relates to a method for production of a naturally folded eukaryotic protein containing two or more cysteines linked by disulfide bridges. The method comprises co-expression and secretion into the periplasm of a molecular chaperone via an expression vector coding for the chaperone. The expression vector also encodes a signal sequence. The method is useful for producing a naturally folded eukaryotic protein such as an antibody, antibody fragment, interferon, protein hormone or a protease containing two or several cysteines linked by disulfide bridges. The present sequence is a fusion protein composed of the PelB signal sequence and ScFvOxazolon. This sequence was used in the method of the present invention Producing naturally folded eukaryotic proteins e.g. antibodies, interferon, hormones or proteases that contain two or several cysteines linked by disulfide bridges comprises co-expression of a molecular 19; 35pp; English. Disclosure; Page chaperone.

AA; Sequence 255

255 88 7 15 0 Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps: 3.12e-38 444.00 86.36% 80.00% 65.88% Percent Similarity: Best Local Similarity: Query Match: Alignment Scores: Score:

(1-255) $(1-359) \times AAB74199$ US-10-049-868A-1

Qy	Ц	1 GACATTGAGCTCACCCAGTCTCCAGCAATCATGTCTGCATCTCCAGGGGGAGAAGGTCACC 60
Db	132 A	
Qy	61 A	TGACCTGCAGTGCCAGTTCAAGTGTTAATTACATGCACTGGTTCCAGCAGGAGTCGGGC 120
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QQ	172 T	ThrSerProLysArgTrp11eTyrAspThrSerLysLeuSerSerGlyValProAlaArg 191
٥٧	181 C	CTCAGTGGCAGTGGGACTGGGAATTCACCCTGGAAATCAGTAGAGTGAAGGCTGAG 240
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QY	241 G	GATGTGGGTGTGTGTGAGAGTTTGTAGAGTATCCGCTCACGTTCGGTGCTGGG 300
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Search completed: October 13, 2004, 13:21:04 Job time: 84.0429 secs

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Sequence 34, Appl Sequence 2, Appli Sequence 251, App Sequence 243, App Sequence 243, App Sequence 241, App Sequence 241, App Sequence 241, App Sequence 5, Appli Sequence 5, Appli Sequence 6, Appli Sequence 242, App Sequence 242, App Sequence 242, Appli Sequence 242, Appli Sequence 240, Appli Sequence 252, Appli Sequence 252, Appli Sequence 252, Appli Sequence 252, Appli Sequence 256, Appli Sequence 255, Appli Sequence 28, Appli Sequence 26, Appli Sequence 6, Appli
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Sequence 34, Application US/10410907A;
Publication No. US20030215880A1
GENERAL INFORMATION:
APPLICANT: Dennis R. Burton
APPLICANT: Anthony Williamson
APPLICANT: Anthony Williamson
APPLICANT: Gianluca Moroncini
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 22908-1229
CURRENT APPLICATION NUMBER: US/10/410,907A
CURRENT FILING DATE: 2003-04-08
PRIOR APPLICATION NUMBER: 60/371,610
PRIOR APPLICATION NUMBER: 60/371,610
PRIOR FILING DATE: 2002-04-09
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
ILENGTH: 215
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
COTHER INFORMATION: D18 Light Chain
US-10-410-907A-34

US-10-738-809-2

US-10-738-809-2

US-10-803-652-251

US-10-803-653-251

US-10-803-653-241

US-10-803-653-241

US-10-803-653-241

US-10-803-653-241

US-10-803-653-241

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US-10-803-653-240

US-09-976-787-8

US-09-976-787-8

US-10-803-653-240

US-09-965-198-8

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-Q=/Cgn2 1/USPTO spool p/US10049868/runat 13102004 132820 24900/app_query.fasta_
-DB=Published Applications AA -QFMT=fastan -SUFFIX=rapb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
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-MAXLEN=200000000 -USER=US10049868 @CGN 1 1 260 @runat 13102004 132820 24900
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-LONGLOG -DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
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| cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
| cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
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Sequence 2, Application US/10344779

Publication No. US2030211106A1

GENERAL INFORMATION:
APPLICANT: SWEET, RAYMOND W.
APPLICANT: TORNETTA, MARK A.
APPLICANT: TORNETTA, MARK A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES
TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS
FILE REFERENCE: GPS003
CURRENT FILING DATE: 2003-02-17
PRIOR APPLICATION NUMBER: US 60/226,524
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-049-868A-1 (1-359) x US-10-344-779-2 (1-129)

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publication US/10738809

geduence 2, Application US/10738809

publication No. US20040171117A1

general INFORMATION:

APPLICANT: SWEET, RAYMOND W.

APPLICANT: TRUNBET, RAYMOND W.

APPLICANT: TRUNBET, ALENSEGED

APPLICANT: TRUNBH, ALENSEGED

APPLICANT: WATTAM, TREVOR A.

TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLONAL ANTIBODIES

TITLE OF INVENTION: USEFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

FILE REFERENCE: GPS0033

CURRENT FILING DATE: 2003-12-17

PRIOR APPLICATION NUMBER: US/10/344,779

PRIOR APPLICATION NUMBER: US/10/344,779

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR APPLICATION NUMBER: US 60/226,524

PRIOR APPLICATION NUMBER: US 60/230,639

PRIOR PILING DATE: 2000-09-07

NUMBER OF SEQ ID NOS: 18

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2

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GENERAL INCORMINION
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APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: Medical Research Council
APPLICANT: Hogenboom, Hendricus
APPLICANT: Hogenboom, Hendricus
APPLICANT: Griffithis, Andrew
APPLICANT: Griffithis, Andrew
APPLICANT: Griffithis, Andrew
APPLICANT: Clackson, Timothy
APPLICANT: Marter, Gregory
APPLICANT: Marter, Gregory
APPLICANT: Marter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REPREMENT: 1300-0013
CURRENT APPLICATION NUMBER: GB 9015198.6
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PLILNG DATE: 1990-10-12
PRIOR PRILING DATE: 1990-10-12
PRIOR PLILNG DATE: 1991-03-06
PRIOR PLILNG DATE: 1991-03-07
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR PLILNG DATE: 1993-07-07
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| AspAlaAlaThrTyrTyrCysGlnGlnArgSerThrTyrProLeuThrPheGlyAlaGly
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                                                                                                                                                            Sequence 251, Application US/10803622 Publication No. US20040157214A1 GENERAL INFORMATION:
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SEQ ID NO 251
LENGTH: 108
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity
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US-10-803-622-251
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APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to C
TITLE OF INVENTION: Their Therapeutic use i
FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1997-04-29
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
SOFTWARE: Patentin version 3.1
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Sequence 17, Application U
Patent No. US20020142359A1
GENERAL INFORMATION:
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ORGANISM: Artificial
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GENERAL INFORMATION:
APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Mcdical Research Council
APPLICANT: Mcdical Research Council
APPLICANT: Mcdical Research Council
APPLICANT: Hoolego, Anthony
APPLICANT: Hoolego, Kevin
APPLICANT: Hoolego, Kevin
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, Cames
APPLICANT: Hoolego, Timechy
APPLICANT: Griffiths, Marks, James
APPLICANT: Griffiths, Calegory
APPLICANT: Winter, Gregory
APPLICANTON NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR PILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-11-12
PRIOR APPLICANTON NUMBER: GB 90245.4
PRIOR APPLICANTON NUMBER: GB 90246.4
PRIOR APPLICANTON NUMBER: GB 901674.9
PRIOR APPLICANTON NUMBER: GB 901679.4
PRIOR FILING DATE: 1991-05-16
PRIOR FILING DATE: 1991-05-15
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICANTON NUMBER: BCCCOT
NUMBER OF SEQ ID NOS: 272
SOFTWARE: PRIOR PRIOR PATEIN VETFION 3.1
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US-10-803-653-251
; Sequence 251, Application US/10803653
; Publication No. US20040157215A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge Antibody Technol
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ORGANISM: Artificial Sequence
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APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology Limited
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
APPLICANT: Pope, Anthony
APPLICANT: Fope, Anthony
APPLICANT: Johnson, Kevin
APPLICANT: Jackson, Ronald
APPLICANT: Jackson, Ronald
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, James
APPLICANT: Marks, James
APPLICANT: Minter, Timothy
APPLICANT: Minter, Timothy
APPLICANT: Monter, Timothy
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APPLICANT: Monter, Gregory
APPLICANT: Monter, Gegory
APPLICANT: Monter, Gegory
APPLICANT: WINBER: GB 9015198.6
FRIOR APPLICATION NUMBER: GB 9022845.3
FRIOR FILING DATE: 1990-10-19
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1990-10-15
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1990-10-15
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1991-05-15
FRIOR APPLICATION NUMBER: US 977/971,857
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ThrLysLeuGluIleLysArgAla 108
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Sequence 243, Application US/10803622
Publication No. US20040157214A1
GENERAL INFORMATION:
APPLICANT: Cambridge Antibody Techno
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APPLICANT: Chiswell, David
APPLICANT: Winter, Gregory
APPLICANT: Winter, Timothy
TITLE OF INVENTION: Methods for Producing Members of
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
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Cambridge Antibody Technology

Cambridge Antibody Technology Limited

F: Medical Research Council

F: McCafferty, John

F: Pope, Anthony

F: Johnson, Kevin

F: Griffiths, Andrew

F: Griffiths, Andrew

F: Holliger, Rasper

F: Marks, James
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Matches:
Conservative:
Mismatches:
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Publication No. US20040157215A1
GENERAL INFORMATION:
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
SOFTWARE: Patentin version 3.1
SEQ ID NO 243
LENGTH: 108
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                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Query Match:
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Pred. No.:
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US-10-803-653-243
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from mouse immunized with 2-phenyl-5-oxazolone
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Publication No. US20040157214A1;
GENERAL INFORMATION:
APPLICANT: Cambridge Antibody Technology
APPLICANT: Medical Research Council
APPLICANT: McCafferty, John
APPLICANT: McCafferty, John
APPLICANT: Hoogenboom, Hendricus
APPLICANT: AppLICANT: Anthony
APPLICANT: AppLICANT: Andrew
APPLICANT: AppLICANT: Andrew
                                                                                                                                                                                                                                                                                                                                                                                                    Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (1-108)
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR PILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: BCT/GB91/01134
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 243
LENGTH: 108
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-049-868A-1 (1-359) x US-10-803-653-243
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80.56%
66.47%
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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US-10-803-622-241
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Pred. No.:
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; OTHER INFORMA
US-10-803-653-241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: VL of scFv from mouse immunized with US-10-803-622-241
      APPLICANT: Jackson, Konald
APPLICANT: Holliger, Kasper
APPLICANT: Holliger, Kasper
APPLICANT: Clackson, Timothy
APPLICANT: Chiswell, David
APPLICANT: Chiswell, David
APPLICANT: Bonert, Timothy
APPLICANTON NUMBER: US/10/803,622
CURRENT APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-11-12
PRIOR PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-05-15
PRIOR FILING DATE: 1991-05-15
PRIOR FILING DATE: 1991-05-16
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1993-01-08
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Conservative:
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Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 272
SOFTWARE: PatentIn version 3.1
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446.00
87.04%
80.56%
66.17%
Jackson, Rona.
Holliger, Kasy
Marks, James
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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ENGTH: 108
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                                                                                                                           GENERAL INFORMATION:
APPLICANT: Cambridge Antibody Technology
APPLICANT: Cambridge Antibody Technology
APPLICANT: Medical Research Council
APPLICANT: Mcafferty, John
APPLICANT: Mcafferty, John
APPLICANT: Mcafferty, John
APPLICANT: Hoogenboom, Hendricus
APPLICANT: Johnson, Kevin
APPLICANT: Griffiths, Andrew
APPLICANT: Marks, James
APPLICANT: Marks, James
APPLICANT: Minetr, Greegory
APPLICANT: Winter, Greegory
APPLICANT: WINTER: 2090-10-19
FRIOR FILING DATE: 1990-10-19
FRIOR APPLICATION NUMBER: GB 9022845.3
FRIOR FILING DATE: 1990-10-19
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1991-03-06
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1991-03-06
FRIOR FILING DATE: 1991-03-06
FRIOR APPLICATION NUMBER: GB 9104744.9
FRIOR FILING DATE: 1991-03-06
FRIOR APPLICATION NUMBER: GB 910474.9
FRIOR FILING DATE: 1991-03-06
FRIOR FILING DATE: 1993-01-08
FRIOR FILING DATE: 
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Mismatches:
Indels:
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RESULT 10
US-10-803-653-241
; Sequence 241, Application US/10803653
; Publication No. US20040157215A1
; GENERAL INFORMATION:
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Best Local Similarity
Query Match:
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83

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ThrCysSerAlaSerSerValSerTyrMetAsnTrpTyrGlnGlnLysSerGlyThr
                                                                                                                                                                                   CORRESTONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz
STREET: One Liberty Place - 46th Fl
CITY: Philadelphia
STATE: PA
COUNTRY: USA
ZIP: 19103
ZIP: 19103
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/704,352
FILING DATE: 07-Nov-2003
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0057
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/795,515
FILING DATE: 28-FEB-2001
APPLICATION NUMBER: 08/846,658
FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
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LENGTH: 235 amino aci
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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445.00
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SEQUENCE TYPE: protein;
US-10-704-352-5
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Local Similarity
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                                                                             184
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                                                                                                                                                                                        Sequence 5, Application US/09795515
; Publication No. US20030039645A1
; GENERAL INFORMATION:
; APPLICANT: Athwal, Diljeet S.
APPLICANT: Emtage, John S.
; TITLE OF INVENTION: Humanised Antibodies
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & 1
STREET: One Liberty Place - 46th Floor
; CITY: Philadelphia
; STATE: PA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Version #1.25
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,515
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/846,658
FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yacko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-00
                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pelacin
                                                                                                             ACCAAGCTGGAGCTGAAACGGGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
TYPE: amino acid
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ZIP: 19103
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Matches:
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                                                              US-10-704-352-5
; Sequence 5, Application US/10704352
; Publication No. US20040071693A1
; GENERAL INFORMATION:
; APPLICANT: Adair, John R.
Athwal, Diljeet S.
Emtage, John S.
; TITLE OF INVENTION: Humanised Antibodies
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
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Percent Similarity:
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Place - 46th Floor
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COUNTRY: USA
ZIP: 19103

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/704,071

TITING DATE: 07-Nov-2003
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Adair, John R.
Athwal, Diljeet S.
Emtage, John S.
TITLE OF INVENTION: Humanised Antibodies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mac
STREET: One Liberty Place - 46th Floo:
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FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0057
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
  (1-235)
 US-10-704-352-5
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SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID N
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US-09-144-886-82
i Sequence 82, Application US/09144886
j Patent No. US20020155114A1
j GENERAL INFORMATION:
   APPLICANT: Marks, James D
   APPLICANT: NUMBER: US/09/144, 886
   CURRENT FILING DATE: 1998-08-31
   NUMBER OF SEQ ID NOS: 98
   SOFTWARE: PatentIN Ver. 2.0
   SEQ ID NO 82
   LENGTH: 107
   TYPE: PRT
   ORGANISM: Artificial Sequence
   FEATURE:
   CORGANISM: Artificial Sequence
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GENERAL INFORMATION:

APPLICANT: Marks, James D

APPLICANT: Amersdorfer, Peter

ITTLE OF INVENTION: Therapeutic Monoclonal Antibo

TITLE OF INVENTION: Botulinum Neurotoxins

ITTLE OF INVENTION: Botulinum Neurotoxins

CURRENT APPLICATION NUMBER: US/09/144,886

CURRENT FILING DATE: 1998-08-31

NUMBER OF SEQ ID NOS: 98

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 83

LENGTH: 107

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

COTHER INFORMATION: Description of Artificial Sequence

COTHER INFORMATION: 1G7 region VL epitope 1

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US-09-144-886-83
; Sequence 83, Application US/09144886
; Patent No. US20020155114A1
; GENERAL INFORMATION:
; APPLICANT: Marks, James D
; APPLICANT: Amersdorfer, Peter
; TITLE OF INVENTION: Therapeutic Monoclonal Ant
; TITLE OF INVENTION: Botulinum Neurotoxins
; FILE REFERENCE: 2500.117USO
; CURRENT APPLICATION NUMBER: US/09/144,886
; CURRENT FILING DATE: 1998-08-31
; NUMBER OF SEQ ID NOS: 98
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 83
; LENGTH: 107
; TYPE: PRT
; ORGANISM: Artificial Sequence
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C;Date: 08-May-1995 #sequence_revision 03-Aug-1995 #text_c C;Accession: S52446
R;Berdoz, J.; Kraehenbuhl, J.P.
submitted to the EMBL Data Library, November 1994
A;Description: Specific amplification by the polymerase ch A;Reference number: S52445
A;Accession: S52446
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-141 <BER>
A;Cross-references: EMBL:X82692; NID:g67341; PIDN:CAA5801
C;Genetics:
A;Introns: 16/1
C;Superfamily: immunoglobulin V region; immunoglobulin hom C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;34-116/Domain: immunoglobulin hom C;Keywords:
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-Q=/Cgn2 1/USPTO_spool_p/US10049868/runat_13102004_132817_24805/ag
-Q=/Cgn2 1/USPTO_spool_p/US10049868/runat_13102004_1200FCL=0 -LOOPEL-ODEB=PIR 79 -QFMT=fastan -SUFFIX=rpr -MINMATCH=0.1 -LOOPCL=0 -LOOPEL-ODEB=PIR 79 -QFMT=fastan -SUFFIX=rpr -MINMATCH=0.1 -LOOPCL=0 -LOOPEL-ODEB-PIR 79 -QFMT=fastan -SUFFIX=rpr -MINMATCH=0.1 -LOOPCL=0 -MINITEN=0 -MINITEN=0 -ALIGN=15 -MC-OUTFMT=pto -NORM=ext -HEAPSIZE=50 -MINITEN=0 -MAXLEN=2000000000 -OUTFMT=pto -NORM=ext -HEAPSIZE=50 -MINITEN=0 -MAXLEN=20000000000 -OUTFMT=pto -NORM=ext -HEAPSIZE=50 -MINITEN=0 -MAXLEN=2000000000 -OUTFMT=pto -NORM=ext -HEAPSIZE=50 -MINITEN=0 -MAXLEN=2000000000 -OUTFMT=pto -NORM=ext -HEAPSIZE=50 -MAIT -DSPBLOCK=100 -LONGLOG -OUTFMT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0 -DEV_TIMEOUT=120 -WARN TIMEOUT=0.5 -DELOP=6 -DELEXT=7
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23-Jul-1999

#text_change

Oy 12 Db 6 Oy 18		chain V 5: Mus mu 56-Jan-19 10n: S385 1er, M.; 1 to the 1 to the 1 to the 1 to 1 t
Oy 24 Db 10 Oy 29	1 AAATGAACAGTCTGCAGACTGATGACACACCCATGTACTACTGCCAGAGATCG 	atus: preliminary lecule type: mRNA sidues: 1-117 <mon> oss-references: EMBL:X75099; NID:941 perfamily: immunoglobulin V region;</mon>
H		ywords: neterotetramer; ımmur -97/Domain: immunoglobulin ho
Qy 35 Db 14	58 TCCTCA 363 40 SerSer 141	or Scores: 8.14e-45 533.50 Conservati
RESULT 2 S11100		Match: 79.27% Indels:
Ig heavy ch. C;Species: I	region (clone NQ2-20.5.3) - mouse (fragment) usculus (house mouse)	2 (1-1) 633060 × (636 1) 6 k030 080
C;Date: 19-1 C;Accession	997 #sequence_revision 23-Aug-1997 #text_(100 Gwiffiths G M : Markham A F : Milstein	383) X 338383 (I-II) BAGCTGCAGGAGTCTGGACCTGC
Nature 304, A;Title: mR	324, 1983 quences define an unusually restricted IgG	
A; Reference A; Accession	er: S07331; MUID:83271467; PMID:6877353 100	OY 61 ACTTGCACTGTCTCTGGGATTTCATTAAACAGATATGG
A;Molecule A;Residues: C:Superfami	mKNA 2 <nat> mmunoqlobulin V region: i</nat>	ThrCysThrValSerGlyPheSerl
F; 15-97/Dom	immunoglobulin homology <imm></imm>	
Alignment Sco Pred. No.: Score:	res: 4.6e-45 Length: 536.00 Matches:	LeuGluTrpLeuG TCCAGACTGAGCA
Percent Sim Best Local	arity: 87.50% Conservative: 1 milarity: 86.67% Mismatches: 7	
Query match DB:	Gaps:	Oy 241 AAATGAACAGTCTGCAGACTGATGACAGCCATGTP
-10-049-	A-2 (1-363) x S11100 (1-112)	
Qy Dp	1 CAGGIGCAGCIGCAGGAGICIGGACCIGGCGTGGCGCCCTCACAGAGCCTGTCCATC 60	Qy 301 ACTATGATTACGGCCTATGCTATGGACTACTGGGCCCTATGGACTACTGGGCCCATGGCCCATGGACTACTGGGGCCCATGCTATGGACTACTGGGGCCCATGGACTACGGGCCCATGGACTACTGGGGCCCATGGACTATGGGACTACTGGGGCCCATGGACTATGGGACTACTGGGACTACTAGGACTACTACTACTACTACTACTACTACTACTACTACTACTA
Sy da	61 ACTIGCACTGTCTCTGGGATTTCATTAAACAGATATGGTGTACACTGGGTTCGCCAGCCT 120	chain V region (E3) - mou
Οχ 1	121 CCAGGAAAGGGTCTGGAGTGGAGTAATATGGACTGGTGGAAGCACAAATTATAAT 180	iles: Mus musculus 9: 07-Jun-1990 #seq 9ssion: PL0087
qa	uGluTrpLeuGlyValIleTrpAl	c, K.; Hasemann, C.; Pollok, B.; Alkan, S.S. p. Med. 169, 519-533, 1989
Qy 10p	181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAACTCCAAGAGCCAAGTTTTCTTA 240 	le: Structural characte erence number: PL0080; ession: PL0087
Qy 2	41 AAAATGAACAGTCTGCAGACTGATGACACAGCCATGTACTACTGTGCCAGAGATCGATC	scure type: mknA idues: 1-120 <mbe> ss-references: GB:X58593</mbe>
	LysmetAsnSerLeuGlnThrAspAspThrAlaMetTyrTyrCysAlaArgAspArgGly 10	Experimental source: strain BALB/C Note: the sequence shown here is from
Qy 3	301 ACTATGATTACGGCCTATGCTATGGACTACTGGGGCCAAGGGACCACGGTCACCGTCTCC 360 	ramily: immunogiobulin v region; immunogi rds: heterotetramer; immunoglobulin /Domain: immunoglobulin homology <imm></imm>
RESULT 3 S38563		Alignment Scores: Pred. No.: 1.61e-44 Length:

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ollok, B.; Alkan, S.S.; Brait, M.; Slaoui, M.; Urbain, J.; C
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erization of antiidiotypic antibodies; evidence that Ab2s ar
MUID:89094248; PMID:2492056
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ibrary, September 1993
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19 heavy chain V region - mouse (fragment) 19 heavy chain V region - mouse) 19 heavy chain V region - mouse) 19 heavy chain Secons chain Chair	Alignment Scores: Pred: No.: Score: S27.00 Conservative: 102 Secret Similarity: 85.954 Mismatches: 9 Best Local Similarity: 87.304 Mismatches: 9 Db

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C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_cha
C; Accession: Sillo6
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG r
A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Accession: Sillo6
A; Molecule type: mRNA
A; Residues: 1-114 <NAT>
C; Superfamily: immunoglobulin V region; immunoglobulin homol
F; 15-97/Domain: immunoglobulin homology <IMM>
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C; Date: 19-Mar-1997 #sequence_revision 23-Au
C; Accession: S11102
R; Kaartinen, M.; Griffiths, G.M.; Markham, P
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually
A; Reference number: S07331; MUID: 83271467; F
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A; Molecule type: mRNA
A; Residues: 1-116 <NAT;
C; Superfamily: immunog
F;15-97/Domain: immunog
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C; Species: Mus musculus (house mouse)

C; Date: 26-Oct-1989 #sequence_revision 31-Dec-1993 #text_change 23-Jul-1999

C; Accession: A32456

R; Dombrink-Kurtzman, M.A.; Johnson, L.S.; Riordan, G.S.; Bedzyk, W.D.; Voss
J. Biol. Chem. 264, 4513-4522, 1989

A; Title: Variable region primary structures of a high affinity anti-fluoresc
A; Reference number: A32456

A; Reference number: A32456

A; Status: preliminary

A; Molecule type: mRNA
A; Residues: 1-139 < DOM>
A; Cross-references: GB: J04609; NID: g556316; PIDN: AAA50298.1; PID: g556317

A; Note: the authors translated the codon CAC for residue 20 as Gln, and CAC
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 34-116/Domain: immunoglobulin homology < INM>
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Matches:
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∑ . ⊡ residue

Scores:

Alignment

C; Accession: S11103 R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C. Nature 304, 320-324, 1983 A; Title: mRNA sequences define an unusually restricted IgG response to 2-phenyloxazolon A; Reference number: S07331; MUID:83271467; PMID:6877353 A; Accession: S1313 A; Molecule type: mRNA A; Residues: 1-115 <nnt> C; Superfamily: immunoglobulin V region; immunoglobulin homology F; 15-97/Domain: immunoglobulin homology <imm> Alignment Scores: Score: 3.51e-43 Length: 101 Pred. No.: 517.00 Matches: 101 Percent Similarity: 85.12% Mismatches: 10 Query Match: 2 Best Local Similarity: 83.47% Mismatches: 1 DB: Gabs: 1</imm></nnt>	US-10-049-868A-2 (1-363) x S11103 (1-115) QY 1 CAGGTGCAGCTGCAGGAGTCTGGACCTGGCCTGGTGGCGC	81 LysMet 301 ACTATG 101 Ala 361 TCA 36 113 Ala 11		int Scores: 5.54e-43 10.: 515.00 Similarity: 84.30% Cal Similarity: 82.64% Match: 76.52% G
	Fasour 10 State 109 Ig heavy chain V region (clone NQ5-78.2.6) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-Jan-2000 C;Accession: S11108 R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C. R;Kartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C. Nature 304, 320-324, 1983 A;Title: mRNA sequences define an unusually restricted IgG response to 2-phenyloxazolone A;Reference number: S07331; MUID:83271467; PMID:6877353 A;Recession: S11108 A;Residues: 1-112 <nat> C;Superfamily: immunoglobulin V region; immunoglobulin homology A;Residues: 1-112 <nat> C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-112 <nat> C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-112 <nat> C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-112 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-12 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-13 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-13 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-13 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-12 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-13 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-13 Sores: C;Superfamily: immunoglobulin homology <inm> A;Residues: 1-146-78 Sores: C;Superfamily: immunoglobulin homology <inm 1-146-78="" a;residues:="" c;superfami<="" sores:="" td=""><td><pre>th:</pre></td><td>21 ThrcysthrvalSerGly***SerLeuThrSerTyrGlyvalHisTrpvalArgGl 121 CCAGGAAAGGTCTGAGTGGTGGTAATATGGACTGGTGGAAGCACAAATTA 121 CCAGGAAAGGTCTGAGTGGTGGTGGAGTTATAGGACTGGTGGAAGCACAAATTA 11 ProGlyLysGlyLeuGluTrpLeuGlyVallleTrpAlaGlyGlySerThrAsnTy 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACACTCCAAGAGCCAAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGCCAAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCGAAAATTAAAAATGAACAGACTGGAGATCGAAAAATGAAAAAAAA</td><td>Oy 301 ACTATGATTACGGCCTATGCTATGGACTACTGGGGCCAAGGGACCACGGTCACCGTCTCC 360 </td></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></inm></nat></inm></nat></inm></nat></nat>	<pre>th:</pre>	21 ThrcysthrvalSerGly***SerLeuThrSerTyrGlyvalHisTrpvalArgGl 121 CCAGGAAAGGTCTGAGTGGTGGTAATATGGACTGGTGGAAGCACAAATTA 121 CCAGGAAAGGTCTGAGTGGTGGTGGAGTTATAGGACTGGTGGAAGCACAAATTA 11 ProGlyLysGlyLeuGluTrpLeuGlyVallleTrpAlaGlyGlySerThrAsnTy 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACACTCCAAGAGCCAAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGCCAAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCCAAGAGTTTT 181 TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACAACTCGAAAATTAAAAATGAACAGACTGGAGATCGAAAAATGAAAAAAAA	Oy 301 ACTATGATTACGGCCTATGCTATGGACTACTGGGGCCAAGGGACCACGGTCACCGTCTCC 360

Db 61 SerAlaLeuLysSerArgLeuSerIleSerLysAspAsnSerLysSerGlnIlePheLeu 80 Oy 241 AAAATGAACAGTCTGCAGACTGATGACACAGCCATGTACTGTGCCAGAGATCGATC	Adding the area chain vegion (anti-idictypic) - mouse clapscales has macking the content of the	(
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Ig heavy chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1998 #sequence_revision 19-Mar-1998 #text_change 21-C;Accession: S26322
R;Accession: S26322
R;Stark, S.E.; Caton, A.J.
J. Exp. Med. 174, 613-624, 1991
A;Title: Antibodies that are specific for a single amino acid inter A;Reference number: S26309; MUID:91341421; PMID:1908510
A;Accession: S26322
A;Molecule type: mRNA
A;Residues: 1-106 <STA>
A;Cross-references: EMBL:X59182
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;9-91/Domain: immunoglobulin homology <IMM>
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STRAIN=FVB/N; TISSUE=Colon;

X MEDLINE=22388257; PubMed=12477932;

A Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

B Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

B Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

A Jones S.J., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human and initial analysis of more than 15,000 full-length human
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01-DEC-2001 (TrEMBLrel. 19, La
01-MAR-2004 (TrEMBLrel. 26, La
1gh-VJ558 protein.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Strausberg R.;
       Proc. Natl. Acad.
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[1]
SEQUENCE FROM N.A.
STRAIN=BALB/C;
MEDLINE=98169018; PubMed=9510199;
Hawlisch H., Frank R., Hennecke M., Baensch M., Sohns B., Bautsch W., Kola A., Klos A., Koehl J.;
"Site-Directed C3a-Receptor Antibodies from Phage Display
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  databases
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Matches:
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  EMBL/GenBank/DDBJ
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Submitted (JUL-2001) to the EMBL/Gen EMBL; BC011181; AAH11181.1; -. PIR; F33932; F33932.
HSSP; P01820; 1A7N.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; C1-set; Z.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN 2.
SEQUENCE 482 AA; 51864 MW; 312E0
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Rodentia;
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01-JUN-2001 (TrEMBLrel. 17, L)
01-OCT-2003 (TrEMBLrel. 25, L)
Single chain Fv (Fragment).
Mus musculus (Mouse).
Plasmid phEN1.
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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P01819;

21-JUL-1986 (Rel. 01, Last sequence update)

21-JUL-1986 (Rel. 01, Last sequence update)

29-MAR-2004 (Rel. 43, Last annotation update)

Ig heavy chain V region MOPC 141 precursor.

Mus musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mucbi TaxID=10090;
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MEDLINE=81012133; PubMed=6774258;
Sakano H., Maki R., Kurosawa Y., Roeder W., Tonegawa S.;
Sakano H., Maki R., Kurosawa Y., Roeder W., Tonegawa S.;
"Two types of somatic recombination are necessary for the of complete immunoglobulin heavy-chain genes.";
Nature 286:676-683(1980).
-!- MISCELLANEOUS: The sequence shown is translated from differentiated gene isolated from a myeloma that seczel. SIMILARITY: Contains 1 immunoglobulin-like domain.
                                                                                                                                                     CRC64
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Mismatches:
Indels:
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Matches:
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J. Immunol. 160:2947-2958(1998).

EMBL; AJ222590; CAA10890.1; -.

PIR; F33932; F33932.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig-v.

Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

Plasmid.

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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mu
   This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMB the European Bioinformatics Institute. There are no restriuse by non-profit institutions as long as its content imodified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sior send an email to license@isb-sib.ch).
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21-JUL-1986 (Rel. 01, Created)

21-JUL-1986 (Rel. 01, Last sequence update)

01-OCT-2004 (Rel. 45, Last annotation updat

Ig heavy chain V region PJ14 precursor.

Mus musculus (Mouse).
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Matches:
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                                                          EMBL; J00491; AAA38121.1; -. PIR; A02094; G2MS14.
HSSP; P01820; 1G7J.
InterPro; IPR007110; Ig-like InterPro; IPR003596; Ig_v.
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81.25%
75.78%
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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21-JUL-1986 (Rel. 01, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Ig heavy chain V region MC101 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi.
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=82075900; PubMed=6273429;
Kataoka T., Nikaido T., Miyata T., Moriwaki K., Honjo T.;
"The nucleotide sequences of rearranged and germline immun genes of a mouse myeloma MC101 and evolution of VH genes i J. Biol. Chem. 257:277-285(1982).
-!- SIMILARITY: Contains 1 immunoglobulin-like domain.
          ThrCysThrValSerGlyPheSerLeuThrGlyTyrGlyValAsnTrpV
                                            region MC101
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Best Local Similarity
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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MEDLINE=88176921; PubMed=2451244;
Schwager J., Mikoryak C.A., Steiner L.A.;
Schwager J., Mikoryak C.A., Steiner L.A.;
"Amino acid sequence of heavy chain from Xenopus laevis IgM deduced from cDNA sequence: implications for evolution of immunoglobulin domains.";
Proc. Natl. Acad. Sci. U.S.A. 85:2245-2249(1988).
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01-FEB-1991 (Rel. 17, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Ig heavy chain V region XIG14 precursor (Fragment).
Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
NCBI_TaxID=8355;
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PIR; B31933; B31933.
HSSP; P01820; 1A7N.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin V region; Signa NON_TER
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|ValSerGlyPheGluLeuSerSerTyrHisMetHisTrpIleArgGlnProProGlyLys
CTGCAGGAGTCTGGACCTGGCTGGCGCCCTCACAGAGCCTGTCCATCACTTGCACT
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                              GTCTCTGGGATTTCATTAAACAGATATGGTGTACACTGGGTTCGCCAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; BC073766; AAH73766.1; -.

InterPro; IPR003599; Ig.

InterPro; IPR003597; Ig.cl.

InterPro; IPR003006; Ig.MHC.

InterPro; IPR003596; Ig.v.

InterPro; IPR003596; Ig.v.
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Last annotation update)
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05-JUL-2004 (TrEMBLrel. 2
Hypothetical protein.
Homo sapiens (Human).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus
NCBI_TaxID=9838;
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TISSUE=Peripheral blood;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies reflecting gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ da EMBL; AB091893; BAD00255.1; -.
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Pfam; PF00047; ig; 4.

SMART; SM00409; IG; 2.

SMART; SM00406; IGc1; 3.

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PROSITE; PS50835; IG_LIKE; 4

PROSITE; PS00290; IG_MHC; UN

Hypothetical protein.

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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
(Fragment).
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                 Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N.,
Young D.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fetus.";
Clin. Immunol. Immunopathol. 87:184-192(1998)
EMBL; AF035041; AAD56277.1; ...
PIR; PH0876; PH0876.
PIR; S12416; S12416.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Myosin-reactive autoantibodies in rheumatic
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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73.64%
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MEDLINE=79124695; PubMed=420800;
Chiu Y.-Y.H., Lopez de Castro J.A., Poljak R.J.;
Chiu Y.-Y.H., Lopez de Castro J.A., Poljak R.J.;
"Amino acid sequence of the VH region of human myeloma cryoimmunoglobulin IgG Hil.";
Biochemistry 18:553-560(1979).
-!- MISCELLANEOUS: This chain was isolated from an IgG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CRC64
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P01771;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Ig heavy chain V-III region HIL.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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-!- SIMILARITY: Contains 1 immunoglobulin-like do:
PIR; A02054; G1HUHL.

HSSP; P01772; 2FB4.

GO; GO:0005576; C:extracellular; NAS.

GO; GO:0006955; P:immune response; NAS.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.
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Pyrrolidone
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61.98%
55.87%
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ne P.H.,
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C TISSUE=Primary B-Cells,

X MEDLINE=22388257; PubMed=12477932;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Rahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.

Whiting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.

Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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llia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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"Generation and initial analysis of more than 15,000 full-le
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01-MAR-2004 (TrEMBLrel
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Q6GMX1;
Q5-JUL-2004 (TrEMBLrel. 27, Created)
O5-JUL-2004 (TrEMBLrel. 27, Last sequence update)
O5-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein.
Hypothetical protein.
Homo sapiens (Human).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Strausberg R.;
Strausberg R.;
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ datak
EMBL; BC011857; AAH11857.2; -.
FIR; S15590; S15590.
HSSP; P01820; 1G7J.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGV; I.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN 3.
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                                                   TISSUE FROM THE PARTY PROBLEM OF THE PERCENCE FROM THE SECONDER OF THE SCHOOL OF THE SCHOO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Strausberg R.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; BC073773; AAH73773.1; -.

InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig-1ike.
InterPro; IPR003596; Ig-2.

InterPro; IPR003596; Ig-4.

Pfam; PF07654; C1-set; 3.

SMART; SM00409; IG; 2.

SMART; SM00406; IG-2.

SMART; S
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78
14
27
10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA sequences.";
L. Acad. Sci. U.S.A. 99:16899-16903(2002)
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Matches:
Conservative:
Mismatches:
Indels:
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366.00
71.32%
60.47%
54.38%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-049-868A-2 (1-363)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and mouse cDNA sec
Proc. Natl. Acad.
                                            N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similarity
TaxID=9606;
                                            FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alignment Scores:
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X Figure From N.A.

TISSUE=Primary B-Cells;

A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

B Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

B Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Petres G.J., Abramson R.D., Mullahy S.J.,

B Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

A Villalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Jones S.J., Marra M.A.;

"T. "Generation and initial analysis of more than 15,000 full-length human

"T. "Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              full-length human
                                                                                         CGATCTACTATGATTACGGCCTATGCTATGGACTACTGGGGGC
                                                                                                                          AsnileTrpGly
Hômo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the EMBL/GenBank/DDBJ databases
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29
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Last annotation update)
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Mismatches:
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Matches:
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                                                                                                                                                                                                          CAAGGGACCACGGTCACCGTCTCA
                                                                                                                                                                                                                                    Created)
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Strausberg R.;
Submitted (JUN-2004) to the EMEMBL; BC073765; AAH73765.1; -.
InterPro; IPR003599; IG.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_C1.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_V.
Pfam; PF07654; C1-set; Z.
Pfam; PF07654; C1-set; Z.
SMART; SM00409; IG; 4.
SMART; SM00409; IG; 4.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
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73.55%
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RESULT 15
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ITSSUE=Primary B-Cells;

X MEDLINE=22386257; PubMed=12477932;

X Altschul S.F., Feingold E.A., Grouse L.H., Derge J.G.,

X Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

X Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

X Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

X Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

X Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

X Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

X Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

X Robar S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

X Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

X Nilalon D.K., Muzny D.M., Soderien E.J., Lu X., Gibbs R.A.,

X Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

X Hiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

X Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

X Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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ArgLeuAsnSerValThrAlaAlaAspThrAlaValTyrTyrCysAlaHisGlySerSer
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                                                                                                  ACTIGCACTGTCTCTGGGATTTCATTAAACAGATATGGTGTACACTGGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequences.";
d. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                Last sequence update)
Last annotation update)
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Indels:
Gaps:
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Jones S.J., Marra M.A.;
"Generation and initial analysis
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SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                              363
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05-JUL-2004
05-JUL-2004
05-JUL-2004
                                US-10-049-868A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=IGHD;
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Primates; Catarrhini; Hominidae; Homo.
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TISSUE=Primary B-Cells;
MEDLINE=22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., S
                                                                                                                                                                     CRC64
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76
114
229
3
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Last sequence update)
Last annotation update)
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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                EMBL/GenBank/DDBJ
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UNKNOWN 2.
MW; FBB97C
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Strausberg R.;
Submitted (DEC-2003) to the EMBE BC063384; AAH63384.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig.cl.
InterPro; IPR003596; Ig.MHC.
InterPro; IPR003596; Ig.MHC.
InterPro; IPR003596; Ig.
Pfam; PF07654; Cl-set; 3.
Pfam; PF00047; ig; 4.
SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKN(SEQUENCE S76 AA; 63363 MW;
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Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
"Generation and initial analysis of more than 15,000 full-length human
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Strausberg R.;
Submitted (DEC-2003) to the EMBL/GenBank/DDBJ databases
EMBL; BC063384; AAH63384.1; -.
Hypothetical protein.
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-Q=/cgn2_1/USPTO spool_p/US10049868/runat_13102004_132816_24785/app_query.fasta_1.1038
-Q=/cgn2_1/USPTO spool_p/US10049868/runat_13102004_132816_24785/app_query.fasta_1.1038
-DE=A_Geneseq_23Sep04_-QFMT=fastan -SUFFIX=rag -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
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-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPS1ZE=500 -MINLEN=0 -MAXLEN=200000000
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The VH sequence is one of eight (AAR21264-71) found to be expressed from a single chain Fv library from an immunised mouse. The library produces a diverse repertoire of antibody fragments specific for 2-phenyl-5-coxazolone (phOx). It was prepd. using cDNA generated from mRNA from mice immunised with phOx coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by PCR (see AAQ23474-84) and ligated into fdCAT2 (see AAQ23463) for expression on the phage surface as fusions with gene III. The resulting library of clones was diverse. Twenty three hapten binding clones were sequenced revealing the eight different VH genes (A-H) in a variety of pairings with seven diffrent VK genes (a-g) (see AAR21264-92). Nearly all the VH genes belonged to gp 1, with only one, "E", being of gp 2 (VHOx1). Of the twenty three clones sequenced, only one was of type "E". Most of the clones were VK-d combinations. The Kd of VH-B/VK-d for phOx-GABA was 10 nM. Only two other combinations (of eleven tested) were found to have higher values. This suggests that phage bearing scFv fragments having weak affinities can be selected with antigen, probably due to the avidity of the multiple antibody heads on the phage. See also AAR21260-307, 309-311; AAR22450, 565-581
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                                                                               Hoogenboom HRJ,
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Matches:
Conservative:
Mismatches:
Indels:
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                                        ANTIBODY TECHNOLOGY
                                                                              KS,
                                                                                                                                                                                                 Example 21; Fig 24; 209pp; English.
                                                                              Johnson
Marks
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                                                                                                                                             Producing members of specific recombinant host cells with a
91GB-00004744.
91GB-00010549.
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Holliger
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15-MAY-1991;
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Jackson RH,
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120
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Vascular Adhesion Protein-1 (VAP-1) antibody. This sequence is included
in the specification for comparison with the antibody of the invention.
The murine antibody sequences of the invention may be used in the
production of a chimeric mouse-human anti-VAP-1 antibody. The nucleic
acid molecules, polypeptides or antibodies are useful in treating VAP-1
mediated inflammatory disorders, such as rheumatoid arthritis,
inflammatory bowel disease, autoimmune diseases or psoriasis. The
chimeric VAP-1 antibody is further used for in vitro and in vivo
diagnostic applications, including in vivo immunoscintigraphic imaging of
inflammation sites. The chimeric MAD's of the invention have improved
kinetic properties compared to the corresponding murine antibodies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New chimeric anti-Vascular Adhesion Protein-1 monoclonal antibodies and encoding nucleic acid molecules, useful for diagnosing and treating chronic inflammatory disorders, e.g. rheumatoid arthritis or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACTIGCACTGTCTCTGGGATTTCATTAAACAGATATGGTGTACACTGGGTTCGCCAGCCT
                                                                                                                                                                                          chain;
                                                                                                                                                                                         light;
                                                                                                                                                                          complementarity determining region; CDR; mouse; Vascular Adhesion Protein-1; VAP-1; antibody; heavy; light chimeric; inflammatory disorder; rheumatoid arthritis; inflammatory disease; autoimmune disease; psoriasis;
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Mismatches:
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Best Local Similarity
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                                                                                                                                                  Anti-VAP-1
                                                                                               ADF77174;
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The present invention describes a method (M1) for ablating or killing an aberrant prostate specific membrane antigen (PSMA)-expressing cell (e.g. an epidermal and a dermal cell). M1 comprises contacting the cell, or a vascular endothelial cell proximate to the cell, with an antibody (or its antigen-binding fragment), which binds specifically to the extracellular domain of PSMA in an amount sufficient to ablate or kill the cell. The antibodies have antipsoriatic, antiarthritic, dermatological, cytostatic, antinflammatory and antiallergic activities, and can be used in vaccines. M1 is useful for treating a skin disorder in a subject, by administering to the subject, an amount of an antibody which binds
                                                                                                            300
                                                                                                                                                                  360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ablating/killing aberrant prostate specific membrane antigen-expressing cells for treating skin disorders, by contacting the cell with an antibody that binds to the extracellular domain of prostate specific
                                                  AAGTTTTCTTA
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|nValPheLeu
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                                                                   administering to the subject, an amount of an antibody specifically to the extracellular domain of PSMA (the
                                                                                                                                                                                                                                                                                                                                                                                      sequence MuVHIB
                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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mammal, preferably human and is having, or at risk of, a skin disorder).

The skin disorder is a dermal or an epidermal disorder, and is selected from psoriasis (preferably chronic stationary psoriasis, psoriasis psoriasis, eruptive (gluttate) psoriasis, psoriatic erythroderma, generalised pustular psoriasis, psoriatic erythroderma, cerfoliative dermatitis, pityriasis rubra pilaris, pityriasis rosacea, psoriasis, pityriasis lichenoiders, lichen planus, lichen nitidus, ichthyosiform dermatosis, keratodermas, dermatosis, and prokeratosis, preferably psoriasis. MI is useful for treating a skin disorder such as an inflammatory or neoplastic disorder of the epidermis or dermis, preferably an epidermal precancerous or cancerous lesion. MI is also useful to treat or prevent disorders involving aberrant activity of PSMA-expressing cell, e.g. kidney, liver or brain cell. ACC69816 to ACC69837 and ABR44613 to ABR44733 represent sequences used in the exemplification
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constructed by linking DNA encoding the Scrv (itself generated by linking DNA encoding the morphisms protein seminal plasma protein. The single-chain antibody (Scrv)/human carboxypeptidase A (hCPA) fusion protein (ABP97414), a prodrug of methotrexate-alpha-peptide and a pharmaceutically-acceptable auxiliary for medication. The invention also encompasses the anti-human seminal plasma protein Scrv/hCPA fusion protein, and the nucleic acid encoding it (ABZ75116). This was constructed by linking DNA encoding the Scrv (itself generated by linking DNA encoding the Scrv (itself generated by linking DNA encoding the Scrv (itself generated by linking DNA encoding the sortein monoclonal antibody E4B7) to DNAs encoding human carboxypeptidase A via a linker. The Scrv portion of the fusion protein enables it to be specifically targeted to prostate cancer the hCPA portion can then activate the prodrug which in turn has cytostatic activity towards the cancer cells. The invention permits the fusion protein to be easily produced in large quantities for use in ADEPT for prostate cancer treatment. The present sequence
                                                                                                                                                                                                                                                       (light
                                                                                                                                         prostate cancer
                                                                                                               chain
E4B7 VH
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The kit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical kits for use in the treatment of prostate cancer, co anti-human seminal plasma protein single-chain antibody/human carboxypeptidase fusion protein and a prodrug of methotrexate-alpha peptides.
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                                                                                                               single
ntibody
                                                                                                                                                                                                                                                        E4B7
                                                                                                                                                                                     E4B7
                                                                                                                                                                                     antibody
                                                                                                                                                                                                                                                        antibody
                                                                                     /label= E4B7_derived_SCFv
/note= "Anti-human seminal plasma protein
antibody derived from murine monoclonal an
and VL regions; targets fusion protein to
cells"
                                                                                                                                                                                                                                                                                                                                                                                        carboxypeptidase A; activate prodrug"
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                                                                                                                                                                                 /note= "Corresponds to monoclonal chain variable region" 123. .137
                                                                                                                                                                                                                                                         monoclonal
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                                                                                                                                                                                                                                                                                                              245. .251
/label= ScFv/hCPA_linker
                                                                                                                                                                                                                                                                                                'note= "Encoded by GAC"
                                                                                                                                                                                                                                                                                                                                                       note= "Encoded by AGC"
                                                                                                                                                                                                                                                         to
                                                                                                                                                                                                                                                                       chain variable region"
                                                                                                                                                                                                                                                         "Corresponds
                                                                      Location/Qualifiers
                                                                                                                                                                                                                             ScFv_linker
                                                                                                                                                                                                                                                                                                                                                                               /label= hCPA
/note= "Human alpha-peptide g
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/note= "Co
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/label=
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N-PSDB; ABZ75116.
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(HAOX/) HAO X.
             Homo sapiens
Synthetic.
  musculus
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                                         Chimeric.
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                                                                      Key
Protein
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Human; NKG2D; NKG2D receptor complex; cancer; infectious disease; tumour; autoimmune disease; head; neck; stomach; oesophagus; colon; liver; ovary; intrahepatic bile duct; pancreas; lung; larynx; breast; uterus; cervix; prostate; kidney; testis; thyroid; bladder; brain; melanoma; myeloma; Fv; sarcoma; leukaemia; lymphoma; virus; bacterium; fungus; protozoan; DAP10; helminth; cytostatic; antimicrobial; immunomodulatory; 11B2D10; 6H7E7; 8G7C10; 6E5A7; 11B2D10x4-7; 8G7C10x4-7; 6E5A7x4-7; P4-2; P4-3; P4-14; P5-15; P5-2; P5-3; P5-9; P5-10; P5-11; P5-23; 3B10xP4-3; 3B10xP4-14; P5-3 tetramerisation domain; 3B10xP5-2; 3B10xP5-23.
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Mismatches:
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                      1.46e-4
547.00
91.74%
86.78%
81.28%
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AA;
                                       Percent Similarity:
Best Local Similarity
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DB:
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Pred. No.:
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The invention relates to a multifunctional polypeptide comprising a domain with a binding site that specifically recognises an extracellular group of the NKG2D receptor complex and a second domain which functions as a receptor or ligand. The polypeptide and its associated polynucleotide are used for the preparation of a pharmaceutical composition for the treatment of cancer, infections and/or autoimmune conditions. The cancer may be a tumour of the head and neck, stomach, esophagus, colon, liver, intrahepatic bile ducts, pancreas, lung, larynx, breast, ovary, uterus, cervix, prostate, kidney, testis, thyroid, larynx, breast, ovary, uterus, cervix, prostate, kidney, testis, thyroid, clarynx, breast, ovary, uterus, cavix, prostate, kidney, testis, thyroid, larynx, breast, ovary, uterus, cavix, prostate, kidney, testis, thyroid, clarynx, breast, ovary, uterus, cavix, prostate, kidney, testis, thyroid, larynx, breases can be caused by viruses, bacteria, fungi, protozoa or helminths. The autoimmune diseases include multiple sclerosis, Grave's disease, ankylosing spondylitis, acute anterior uveitis, Goodpasture's syndrome, myasthenia gravis, insulin-dependent diabetes mellitus, rheumatoid arthritis, pemphigus vulgaris and autoimmune hepatitis. Sequences AAU72820-AAU72875 represent the NKG2D receptor and the polypeptides of the invention
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rValThrVal
                                                        nuttiunctional polypeptides comprising binding sites that sprecognize extracellular groups of the NKG2D receptor complex thich function as receptors or ligands, useful for treating confectious diseases.
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Matches:
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Hofmeister
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N-PSDB; AAS97141.
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serSer
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infectious
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The present sequence represents the heavy chain variable region of murine monoclonal antibody 2C4. This antibody binds to human sialoadhesin factor -2 (SAF-2). The antibody is useful for treating or preventing allergic rhinitis, allergies, asthma, anemia, eczema or diseases such as lymphoma, leukemia or systemic mastocytosis in a mammal. It is also useful for detecting the presence of a cell, especially eosinophil in a sample, by detecting binding of the antibody to SAF-2. The antibody can be coupled to toxins, antiproliferative drugs or radionuclides to kill cells in a reas of excessive SAF-2 expression
                                                                       sialoadhesin factor-2; SAF-2; allergic rhinitis;
eczema; lymphoma; systemic mastocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      180
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                                                                                                                                                                                                                                                                                                                                     Novel monoclonal antibody specific for human sialoadhesin factor-2 for diagnosis, prevention, treatment of allergy, asthma, eczema or diseases such as lymphoma, leukemia or systemic mastocytosis, in a mammal.
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Matches:
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                                                                       al antibody 2C4;
asthma; anemia;
eosinophil.
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N-PSDB; AAH78183.
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                                                                       Monoclonal
allergy; as
leukemia; e
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Schleimer F
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 regions.
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Region
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                                                        360
                    300
                                                                            119
                                                                                                                                                                                                                                        Primer; human; immunoglobulin; Ig; variable region; VH; VL; Ck; JH; lymphocyte; vector; soluble; antibody; phage; linker; back; VH3; nested; in-cell PCR; cloning; polymorphic; TCR V; antiphenyloxazolone; hybridoma NQ2/12.4; NQ10/12.5.
 80
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                                                        SerAlaLeuMetSerArgLeuSerIleSerLysAspAsnSerLysSerGlnValPheLeu
                    GAGATCGATCT
                hybridomas - to link together sequences to facilitate analy
                                                                                                                                                                                                                        VH NQ2/12.4-Vk NQ10/12.5 linked peptide sequences #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Winter GP;
                                                                                                                                                                                                                                                                                                                                     114. .115
/note= "Linker peptide"
116. .222
/label= Vkappa_NQ2/12.4
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non-contiguous DNA
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/label= VH_NQ2/12.4
114. .115
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                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                       standard; protein; 222 AA.
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92GB-00012419
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(first en
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N-PSDB; AAQ37462.
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11-JUN-1992;
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19-JUN-1993
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Primer; human; immunoglobulin; Ig; variable region; VH; VL; Ck; JH; lymphocyte; vector; soluble; antibody; phage; linker; back; VH3; nested; in-cell PCR; cloning; polymorphic; TCR V; antiphenyloxazolone; hybridoma NQ2/12.4; NQ10/12.5.
                                                                                                             222
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                                             Conservative:
Mismatches:
                           Length:
Matches:
                                                                 Indels
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128. .235
/label= Vkanna NO3/10 :
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abel= Vkappa_NQ2/12
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538.00
87.60%
85.95%
79.94%
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/label= '
114. .12
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(first en
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19-JUN-1993
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92WO-GB001483

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The sequences given in AAR32840-43 show the mature heavy chain VH domains and the Vk light chain genes of the antiphenyloxazolone hybridomas NQ2/12.4 and NQ10/12.5 which have been linked via a linker peptide by incell PCR. The cDNA encoding these peptides was synthesised using forward primers annealing to the Ck gene and the JH segment, followed by assembly with linker primers, VH back primers based on the VH3 leader sequence and a forward Ck primer nested in respect to the primer used for cDNA. The assembled product within the cells is then amplified with nested primers annealing to the 5' end of the VH gene and the 3' end of the Jk segment. In-cell PCR may be used to determine gene linkage analysis, particularly for the cloning of gene combinations that are polymorphic within a population of cells, such as the rearranged genes for Ig or TCR V regions. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                        analysis
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facilita
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                                                                                                                              to t
                                                                                                                            ell populations, partic. hybridomas more non-contiguous DNA sequences
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Matches:
Conservative:
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92GB-00012419
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N-PSDB; AAQ37459.
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The invention relates to an antibody-toxic group conjugate comprising an antibody that specifically recognises a molecule expressed only on activated T cells, and a toxic group. The T cell molecule is preferably human cytotoxic T lymphocyte associated antigen 4 (CTLA4). The antibody of the invention is a humanised anti-CTLA4 antibody comprising a sequence of 128 or 142 amino acids fully defined in the specification. The antibody-toxic group conjugate is useful for modulating the immune cesponse in a subject suffering from a disorder or condition such as autoimmune disorder, immune response to a graft, allergic response or an immune response to a therapeutic protein. The antibody is also useful for T-cell typing, for isolating specific IL-creeptor-bearing cells or fragments of the receptor, for vaccine preparation, and for determining the effectiveness of an agent to down-region of mouse antibody 26. It was used in the construction of the region of mouse antibody of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antibody
a sequence
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                                                                                        -CTLA4;
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r, allergic
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                                                                                      antigen 4; anti-CTLA:
;; vaccine; antibody;
graft rejection;
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                                                                                                                                                                                                                                                                                                                                                                                                Novel antibody-toxic group conjugate comprising an antibody that recognizes a molecule expressed only on activated T cells, usefu modulating immune response for treating autoimmune disorder, all
                                                                                                                                                                                                                                                                                                                    Morris
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107
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107
                                                                                      Mouse, CTLA4; cytotoxic T lymphocyte associated animunosuppressive; immunomodulator; antiallergic; T cell; humanised antibody; autoimmune disorder; grallergy; antibody 26; heavy chain variable region;
                                                                                                                                                                                                                                                                                                                    Gray
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Matches:
Conservative:
Mismatches:
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Tsurushita
                                                                chain
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Local Similarity
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                                                                antibody
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Pred. No.:
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RESULT **AAG665**

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                                                                            99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide(s)
g. melanoma an
                  -GAT
                                                                                                      'alArgGlnPro
                                                                                                                                                                                                                                                                                                       Monoclonal antibody; MAb; 1A7; GD2; immune response; melanoma; neuroblastoma; glioma; soft tissue carcinoma; small cell carcitumor-associated antigen.
                                                                                             AAAATGAACAGTCTGCAGACTGATGACACAGCCATGTACTACTGTGCCAGA
ThrCysThrValSerGlyPheSerLeuThrSerTyrGlyValTyrTrpV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      noclonal antibody 1A7 and related polynucleotide(s) as useful to treat or palliate a GD2-associated disease,
                                                                                                                                                                                                                                                                                                                                                                                  "residues Xaa are unspecified"
                                                                                                                                                                                                                                                                                      consensus sequence.
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                                                                                                                                                                                                                              119 AA
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                                                                                                                                                                                                                                                                                    variable region
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                       chain resembling 1A7 #7
                       Length:
Matches:
Conservative:
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Indels:
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96US-00591196.
96US-00752844.
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Best Local Similarity:
AA;
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16-JAN-1996;
21-NOV-1996;
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7
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The invention relates to the recurrence and/or development of a ganglioside GD2-associated tumour, e.g. melanoma, in an individual which is delayed by administration of an antibody comprising light and heavy chain variable region sequences of the anti-idiotype monoclonal antibody 1A7. The antibody is used for delaying recurrence and/or development of GD2-associated tumour, e.g. melanoma, neuroblastoma, glioma, sarcoma, or small cell lung cancer, in individual, and for treating individual with GD2-associated tumour. The present sequence is a unique peptide region from an antibody sequence resembling the anti-idiotype antibody 1A7 heavy chain.
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   containing
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 y administering antibody contain sequences
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Matches:
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Mouse; monoclonal antibody; 1A7; anti-idiotype antibody; ganglioside GD2; cytostatic; melanoma; neuroblastoma; small cell lung cancer; tumour; antibody; variable region; VH; VL; glioma; soft tissue sarcoma; vaccine. Anti-idiotype antibody VH region consensus sequence 119 protein; entry) (first standard; 119 Ser 8-DEC-2003 ADC35320 119 ADC35320 RESULT 12 ADC35320

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US-10-049-868A-2 (1-363)

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Alignment

Score:

1.87e-4 535.00 88.43% 85.95% 79.49%

Percent Similarity: Best Local Similarity: Query Match:

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Synthetic. Mus sp.

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The invention relates to a monoclonal antibody 1h7 (an anti-idiotype antibody eliciting an anti-GD2(ganglioside) response). Also included are antibody eliciting an anti-GD2(ganglioside) response). Also included are antibody producing cell deposited under ATCC Accession Nov. HB-11786 (or its progeny), a polynucleotide comprising a sequence encoding a polypeptide with immunological activity of 1A7 (where the polypeptide comprises at least 5 consecutive amino acids from a variable region of 1A7), an isolated polynucleotide comprising a region of at least 20 consecutive nucleotides that is capable of forming a stable duplex with a polynucleotide encoding the light or heavy chain variable region of 1A7 under conditions where the region does not form a stable hybrid with a polynucleotide consisting of a variable region encoding sequence appearing as ADC3531 - ADC35370, a host cell comprising the polynucleotide on a fusion polypeptide comprising 1A7 variable regions and a polynucleotide, a fusion polypeptide comprising 1A7 variable regions and a vaccine comprising the antibodies. The antibodies are useful for eliciting an immune response in an individual, and for treating a GD2-associated diseases is a sesciated disease is an individual thas a clinically detectable tumour, and the method is for palliating the GD2-associated disease. 1A7 is an individual and has been treated and is clinically detectable tumour that was previously detectable time of the administering of 1A7, or for reducing the risk of recurrence of a clinically detectable tumour. And the humanised antibody are tumour cell. The present sequence represents a consensus sequence for the tumour cell. The present sequence represents a consensus sequence for the
                                                                                                                                                                                                                                                                                                                                                                that is capable of glycosphingolipid GD2, e.g., melanoma, glioma,
                                                                                                                                                                                                                                                                                                                                                                 1A7,
                                                                                                                                                                                                                                                                                                                                                               Novel anti-idiotypic monoclonal antibody 1A7, recruiting a tumor-specific response against useful for treating a GD2-associated disease
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                                                                                                                                                                                                                                                                                      Chatterjee
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96US-00591196.
99US-00293533.
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16-JAN-1996;
15-APR-1999;
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Best Local Similarity:
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Pred. No.:
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     Heavy chain; variable region; anti-DNA; monoclonal; antibody; 11f8 g
hairpin; diagnosis; inflammatory glomerulonephritis;
systemic lupus erythematosus; screening; treatment; prevention; SLE;
disease; consensus; putative.
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C
                                                                                    TCGGCTCTCATGTCCAGACTGAGCATCAGCAAAGACACTCCAAGAGC
                                        SerAlaLeuLysSerArgLeuSerIleSerLysAspAsnSerLysSer
                                                                                                                                                                                              Anti-DNA antibody 11f8 group heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-DNA antibody which specifically binds DNA hairpin -develop prods. for diagnosis and treatment of disorders, glomerulonephritis or systemic lupus erythematosus.
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The present sequence is the heavy chain variable region of the group 11f8 putative consensus anti-DNA monoclonal antibody (MAb), which has a high affinity for single stranded DNA, low or no affinity for double stranded DNA and specifically binds a DNA hairpin. The MAb can be used diagnose disorders associated with the pathological complexation of DNA, e.g. inflammatory glomerulonephritis and systemic lupus erythematosus. It can also be used to generate reagents to screen for pharmaceutical agents, and treat and/or prevent an above disorder. The sequence was derived by aligning homologous anti-DNA MAb, whose sequences have been published, as well as several MAb of other specificities obtained from a database search
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Gaps:
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The present invention relates to an anti-Death receptor 4 (DR4) antiborantibodies of the invention are useful for inducing apoptosis in mammalian cancer cells such as colon cancer cells and for treating an immune-related disease in a mammal such as arthritis and autoimmune disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel anti-death receptor 4 antibodies useful for treating cancer immune related disorders such as rheumatoid arthritis, sjogren's syndrome, Grave's disease and diabetes mellitus.
                                                                     antibody; apoptosis; cancer;
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                                                                     DR4;
    standard; protein;
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                                     entry)
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                                                                    Anti-Death
                                                                            autoimmune
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Search completed: October 13, 2004, 13:21:06 Job time : 84.9571 secs

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Sequence 218, Apples Sequence 63, Apple Sequence 12, Appli Sequence 12, Appli Sequence 23, Appli Sequence 23, Appli Sequence 23, Appli Sequence 252, Apples Sequence 252, Apples Sequence 252, Apples Sequence 254, Apples Sequence 212, Apples Sequence 31, Apples Sequence 21, Apples Sequence 21, Apples Sequence 21, Apples Sequence 22, Apples Sequence 24, Apples Sequence 25, Apples Sequence 25, Apples Sequence 25, Apples Sequence 26, Apples Sequence 26, Apples Sequence 53, Apples Sequence 63, Apples Sequence 64, Apples Sequence 65, Apples 65, Apples
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US-10-803-622-218

JOSTIC 218, Application US/10803622

Publication No. US20040157214A1

GENERAL INFORMATION:

APPLICANT: Cambridge Antibody Technology

APPLICANT: Cambridge Antibody Technology

APPLICANT: Medical Research Council

APPLICANT: McCafferty, John

APPLICANT: McCafferty, John

APPLICANT: Hope, Anthony

APPLICANT: Hoogenboom, Hendricus

APPLICANT: Griffiths, Andrew

APPLICANT: Griffiths, Andrew

APPLICANT: Griffiths, Andrew

APPLICANT: Griffiths, Andrew

APPLICANT: Clackson, Timothy

APPLICANT: Winter, Gregory

APPLICANT: Winter, Gregory

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APPLICANT: Winter, Gregory

APPLICANT: Winter, Timothy

TITLE OF INVENTION: Methods for Producing Members of

FILE REFERENCE: 13839-00013
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US-10-153-401-16

US-10-660-128-12

US-08-973-518-4

US-09-007-093-4

US-10-232-187-2

US-10-232-187-2

US-10-232-187-2

US-10-232-187-2

US-10-207-655-254

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US-10-10-351-748-31

US-10-194-975-112

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US-09-842-776A-52

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-Q=/cgn2 1/USPTO spool p/US10049868/runat 13102004 132820 24900/app_query.fasta_1.1038
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-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
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CURRENT FILING DATE: 2004-03-18

PRIOR APPLICATION NUMBER: GB 9015198.6

PRIOR FILING DATE: 1990-07-10

PRIOR APPLICATION NUMBER: GB 9022845.3

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-11-12

PRIOR APPLICATION NUMBER: GB 9024503.6

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1991-03-06

PRIOR FILING DATE: 1991-05-15

PRIOR FILING DATE: 1991-05-15

PRIOR FILING DATE: 1991-07-10

PRIOR FILING DATE: 1993-01-08

PRIOR FILING DATE: 1993-01-08

PRIOR FILING DATE: 1993-01-08

PRIOR FILING DATE: 1995-06-07

NUMBER OF SEQ ID NOS: 272

SOFTWARE: PATENTIN VERSION 3.1

SEQ ID NO 218

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US-10-803-653-218
; Sequence 218, Application US/10803653
; Publication No. US20040157215A1
; GENERAL INFORMATION:
; APPLICANT: Cambridge Antibody Techno
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APPLICANT: Cambridge Antibody Technology Laboration APPLICANT: Medical Research Council APPLICANT: McCafferty, John McCafferty, John APPLICANT: Pope, Atthony APPLICANT: Pope, Atthony APPLICANT: Hoogenboom, Hendricus APPLICANT: Griffiths, Andrew APPLICANT: Holliger, Kasper APPLICANT: Holliger, Kasper APPLICANT: Holliger, Kasper APPLICANT: Holliger, Kasper APPLICANT: Claswell, David APPLICANT: Grieswell, David APPLICANT: Grieswell, David APPLICANT: Minter, Gregory APPLICANT: Minter, Gregory APPLICANT: Minter, 13839-00013 CURRENT FILING DATE: 1990-10-19 PRIOR APPLICATION NUMBER: GB 9015198.6 PRIOR APPLICATION NUMBER: GB 9022845.3 PRIOR APPLICATION NUMBER: GB 9024503.6 PRIOR APPLICATION NUMBER: GB 9104744.9 PRIOR FILING DATE: 1991-03-06 PRIOR FILING DATE: 1991-03-06 PRIOR FILING DATE: 1991-07-10 P
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Matches:
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NUMBER OF SEQ ID NOS: 272
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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89.26%
82.76%
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Best Local Similarity
Query Match:
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Pred. No.:
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Query Match:
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US-10-239-656-63

US-10-239-656-63

Sequence 63, Application US/10239656

Publication No. US20040038339A1

GENERAL INFORMATION:

APPLICANT: RIETHMULLER, GERT
APPLICANT: RIETHMULLER, RALF

APPLICANT: RISCHEL, ROMAN

APPLICANT: HORBISE, RALF

APPLICANT: MAYER, MONIER, RALF

APPLICANT: MAYER, MONIER, RAFRIN

APPLICANT: MAYER, MONIER, ROBERT

TITLE OF INVENTION: TO AN EPITOPE OF THE NKG2D RECEPTOR COMPLEX

TITLE OF INVENTION: TO AN EPITOPE OF THE NKG2D RECEPTOR COMPLEX

TITLE OF INVENTION NUMBER: PCT/EP01/03414

FILE REFERENCE: 2001-03-26

CURRENT FILING DATE: 2001-03-26

PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 92

SOFTWARE: PATCHIN VOY: 2.1

SEQ ID NO 63

LENGTH: 253

LENGTH: 253

LENGTH: 253
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US-10-239-656-63
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Matches:
Conservative:
Mismatches:
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88.52%
81.13%
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Best Local Similarity
Query Match:
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Matches:
Conservative:
Mismatches:
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                                                                                    Sequence 4, Application US/09772103
Sequence 4, Application US/09772103
Publication No. US20020039581A1
GENERAL INFORMATION:
APPLICANT: Carreno, Beatriz M.
APPLICANT: Wood, Clive
APPLICANT: Collins, Mary
APPLICANT: Collins, Mary
APPLICANT: Gray, Gary S.
APPLICANT: Gray, Gary S.
APPLICANT: Hinton, Paul
APPLICANT: Hinton, Paul
APPLICANT: Hinton, Paul
APPLICANT: GNN-009CP
CURRENT HING DATE: 2001-01-23
CURRENT FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.0
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84.55%
79.79%
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LENGTH: 142
TYPE: PRT
ORGANISM: Mus musculus
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Foon, Kenneth A.
Chatterjee, Sunil K.
INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE
TREATMENT OF MELANOMA AND SMALL
                                                                                                                                                                                                                                                                                                                  Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10/153,401
FILING DATE: 27-Aug-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/293,533
FILING DATE: 1999-04-15
APPLICATION NUMBER: US 08/372,676
FILING DATE: 1995-01-17
APPLICATION NUMBER: US 08/591,196
FILING DATE: 1996-01-16
ATTORNEY/AGENT INFORMATION:
NAME: Catherine M. Polizzi
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 304142000202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
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Mismatches:
Indels:
Gaps:
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                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94304-1018
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
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Matches:
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ADDRESSE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CA
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RESULT 5
US-10-153-401-16
; Sequence 16, Application US/10153401
; Publication No. US20030114398A1
; GENERAL INFORMATION:
: APPLICANT: Chatterjee, Malaya
Foon, Kenneth A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 : (415) 813-5600
(415) 494-0792
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TELEX: 706141
NMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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APPLICANT: Ashkenazi, Avi
APPLICANT: Chuntharapai, Anan
APPLICANT: Chuntharapai, Anan
APPLICANT: Chuntharapai, Anan
APPLICANT: Chuntharapai, Anan
TITLE OF INVENTION: DR4 Antibodies and Uses;
FILE REFERENCE: P1245R1P2B
CURRENT APPLICATION NUMBER: US/09/584,166
PRIOR APPLICATION NUMBER: US/09/584,166
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1998-01-26
NUMBER OF SEQ ID NOS: 12
LENGTH: 476

LENGTH: 476
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Sequence 12, Application US/10660128
Publication No. US20040120947A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Sequence is
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Best Local Similarity:
Query Match:
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Publication No. US20010014327A1

GENERAL INFORMATION:

APPLICANT: Anand, Naveen N

APPLICANT: Barber, Brian H

APPLICANT: Cates, George A

APPLICANT: Caterini, Judith E

APPLICANT: Klein, Michel H

TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR DELIVERY OF

TITLE OF INVENTION: ANTIGENS TO SELECTED CELLS OF THE IMMUNE
 -CGATCTACTATGATTACGGCCTATGCTATGGACTACTGGGG
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Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/973,51
FILING DATE: 07-APR-1998
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 140 amino acids
                                                                                                                                                                     8.63e-4
531.50
88.43%
85.12%
78.97%
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Barber, Brian H
APPLICANT: Cates, George A
APPLICANT: Caterini, Judith E
APPLICANT: Klein, Michel H
TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR D
TITLE OF INVENTION: ANTIGENS TO SELECTED CELL
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: Suite 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/007,093
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/483,576
FILING DATE:
ATTORNEY/ACTION NUMBER: US 08/483,576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/483,576
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-765
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09007093
Patent No. US20020025315A1
GENERAL INFORMATION:
APPLICANT: Anand, Naveen N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: Suite 701, 330 Uni
CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
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Barber, Brian H
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L, 330 University Avenue
           Length:
Matches:
Conservative:
Mismatches:
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TO SELECTED
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FILING DATE: 05-May-2003
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPACIANT COMPUTER: IBM PC COMPACIONS OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION DATA:
APPLICATION NUMBER: US 10/428,75
FILING DATE: 05-MAY-2003
                                                     Indels
                                                                                   \times US-09-007-093-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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US-10-428-754-4
; Sequence 4, Application US/10428754
; Publication No. US20030228304A1
; GENERAL INFORMATION:
; APPLICANT: Anand, Naveen N
; Barber, Brian H
; Cates, George A
; Caterini, Judith E
; Klein, Michel H
; TITLE OF INVENTION: CHIMERIC AN
ANTIGENS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Stewart, Michael REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & Mc
STREET: Suite 701,
           -46
          8.63e-4
531.50
88.43%
85.12%
78.97%
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                   Score:
Percent Similarity:
Best Local Similarity:
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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US-10-232-187-2
Sequence 2, Application US/10232187
Publication No. US20030092091A1
GENERAL INFORMATION:
APPLICANT: Abrahamson, Julie A.
APPLICANT: Bochner, Bruce
APPLICANT: Erickson-Miller, Connie L.
APPLICANT: Kikly, Kristine K.
APPLICANT: Schleimer, Robert
APPLICANT: Schleimer, Robert
APPLICANT: Nulku, Turkan E.
TITLE OF INVENTION: Sialoadhesin Factor-2 And FILE REFERENCE: GH50042-1
CURRENT APPLICATION NUMBER: US/10/232,187
CURRENT APPLICATION NUMBER: 60/187,595
PRIOR APPLICATION NUMBER: 60/187,595
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2001-03-05
PRIOR FILING DATE: 2001-03-05
PRIOR FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 60/315,943
PRIOR APPLICATION NUMBER: 60/315,943
                                                                                                                           4
REFERENCE/DOCKET NUMBER: 103:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                           S
                                                                                                                            П
                                                                          LENGTH: 140 amino acids
                                                                       TYPE: amino acid

STRANDEDNESS: single

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TOPOLOGY: linear

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SEQUENCE DESCRIPTION: SEQ

US-10-428-754-4
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531.50
88.43%
85.12%
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Best Local Similarity
Query Match:
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                                            . Sequence: Sheavy chain
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Matches:
Conservative:
Mismatches:
Indels:
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                                          Description of Artificial hybridoma 6H7E7 variable
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US-10-144-644-15

Sequence 15, Application US/10144644

Publication No. US20030059429A1

GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Cook, Richard M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant ILS
TITLE OF INVENTION: Treatment of ILS
TITLE OF INVENTION: Treatment of ILS
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                       Sequence
                                                                                                 3.1e-45
526.00
86.78%
85.95%
78.16%
LENGTH: 119
TYPE: PRT
ORGANISM: Artificial S
FEATURE:
OTHER INFORMATION: Des
OTHER INFORMATION: hyb
                                                                                                                        Percent Similarity:
Best Local Similarity
Query Match:
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                                                                                         Scores
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Matches:
Conservative:
Mismatches:
Indels:
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Sequence 23, Application US/10239656

Publication No. US20040038339A1

GENERAL INFORMATION:

APPLICANT: RIETHMULLER, GERT

APPLICANT: LUTTERBUSE, RALE

APPLICANT: LUTTERBUSE, RALE

APPLICANT: HOFMEISTER, ROBERT

TITLE OF INVENTION: MULTIFUNCTIONAL POLYPEPT

TITLE OF INVENTION: TO AN EPITOPE OF THE NK

FILE REFERENCE: 029976/0106

CURRENT APPLICATION NUMBER: US/10/239,656

CURRENT FILING DATE: 2003-03-06

PRIOR PILING DATE: 2001-03-26

PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 92

SOFTWARE: PATENTIN VET: 2.1
                                              4.0
                                                                                                                                                                                            Gaps:
 PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: 60/394,741
PRIOR FILING DATE: 2002-07-10
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 2
LENGTH: 119
                                                                                                                                    9.54e-4
531.00
91.74%
85.95%
78.90%
                                                                              ; TYPE: PRT
; ORGANISM: Homo sapien
US-10-232~187-2
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Ser 119
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Best Local Similari
Query Match:
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APPLICANT:
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US-10-700-740-15
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Mismatches:
Indels:
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 Version
                                                                                                                          APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION: INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5090
INFORMATION FOR SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                        Length:
Matches:
                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
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US-10-700-740-15
; Sequence 15, Application US/10700740
; Publication No. US20040156850A1
; GENERAL INFORMATION:
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523.00
88.43%
82.64%
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amino acid
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; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-10-144-644-15
                                          CLASSIFICATION:
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Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
OF INVENTION: Recombinant IL5 Antagonists Useful in
Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                    Version #1.30
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Mismatches:
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APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                      SmithKline Beecham Corp
O. Box 1539-UW2220
of Prussia
                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/700,740
FILING DATE: 04-Nov-2003
CLASSIFICATION: <Unknown>
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Matches:
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REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
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   TELEPHONE: 610-270-5024
   TELEFAX: 610-270-5090
ORMATION FOR SEQ ID NO: 15:
   SEQUENCE CHARACTERISTICS:
   LENGTH: 119 amino acids
   TYPE: amino acids
                                                                                                                                                                                                                                 STREET: P.O. Box 1539-UW22
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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523.00
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CORRESPONDENCE ADDRESS
ADDRESSEE: SmithK
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                                     ProGlyLysGlyLeuGluTrpLeuGlyValIleTrpAlaSerGlyGlyThrAspTyrAsn
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                       Hession, Catherine A
Whitty, Adrian
TITLE OF INVENTION: COMMON GAMMA CHAIN BLOCKING AGENTS
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Matches:
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Indels:
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APPLICATION NUMBER: US/09/824,286
FILING DATE: 02-Apr-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/017,466
FILING DATE: 10-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan, Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A006 PCT C
TELEPHONE: 617 679-2000
TELEPHONE: 617 679-2838
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTER READABLE FORM:

ZIP: 02142

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

OFTWARE: Patentin Release #1.0, V
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TOPOLOGY: linear
ECULE TYPE: peptide
GMENT TYPE: internal
UENCE DESCRIPTION: SEQ ID NO: 3
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                                                                                                                                                                                                                                                                                           Burkly, Linda C
Benjamin, Christopher
Hession, Catherine A
                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Biogen, Inc.
STREET: 14 Cambridge Center
CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                        S-09-824-286-3
Sequence 3, Application US/09824286
Patent No. US20020028202A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
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Query Match:
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Ser 119
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; SEQUENCE
US-09-824-286-3
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              US-10-207-655-252

US-10-207-655-252

Sequence 252, Application US/10207655

Publication No. US20030118592A1

GENERAL INFORMATION:

APPLICANT: Ledbetter, Martha S.

TITLE OF INVENTION: BINDING DOMAIN-IMMUNOGLOBULIN FUSION PROTEINS

FILE REFERENCE: 390069.401C1

CURRENT APPLICATION NUMBER: US/10/207,655

NUMBER OF SEQ ID NOS: 426

SOFTWARE: Patentin version 3.0

SEQ ID NO 252

LENGTH: 121
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Matches:
Conservative:
Mismatches:
Indels:
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US-09-824-286-3
                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: fusion polypeptide -10-207-655-252
                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
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Pred. No.:
Score:
Percent Similarity:
Best Local Similarity:
Query Match:
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Search completed: October 13, 2004, 13:53:42	Search Tob ti
121 Ser 121	aga
361 TCA 363	QY
101 SerAsnPheHisTyrTyrValMetAspTyrTrpGlyGlnGlyThrSerValThrValSer 120	QQ
301 ACTATGATTACGGCCTATGCTATGGACTACTGGGGCCAAGGGACCACGGTCACCGTCTCC 360	δλ
81 LysMetAsnSerLeuGlnThrAspAspThrAlaArgTyrTyrCysAlaArgAspGlyTyr 100	qu
241 AAAATGAACAGTCTGCAGACTGATGACACACCATGTACTACTGTGCCAGAGATCGATC	δλ

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5.1.6
Compugen Ltd
GenCore version (c) 1993 - 2004
          Copyright
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- protein search, using sw model OM protein

Seconds 2004, 12:48:23 October 13, on: Run

updates/sec ; Search time 24.2958 (without alignments) 471.266 Million cell

US-10-049-868A-3

1 DIELTQSPAIMSASPGEKVT......GTKLELKRADAAPTVS 909 score: Sequence: Title: Perfect :

119

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Gapext BLOSUM62 Gapop 10.0 Scoring table:

0.5

96216763 residues 283416 segs, Searched:

283416 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 2000000000 Minimum DB Maximum DB

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:* H 0 K 4

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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SUMMARIES	 	\$25058	1001	C440	L007	5437	3056	D001	2959	0526	8007	3056	2788	4904	1111	1111	1112	3056	T040	T040	1111	1111	T039	C607	T040	2959	1111	3245	T040	8007
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g kappa chain V	light c	g kappa chain V	g light chain V	g kappa chain pr	g light chain v	g kappa chain V	g light chain V	g light chain V	g kappa chain V	g kappa chain V	g kappa chain pr	g kappa chain v	g light chain V	g kappa chain pr	g light chain v
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85 6	384	8	8	∞	œ				M	m	376	m	ιų	ď	M

ALIGNMENTS

RESULT

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S25058	
Ig kappa chain	in - mouse
C;Species: N	muscu
C; Date: 19-Mar	-1997 #sequenc
C;Accession:	25058
R; Fischer, F	.; Nie
submitted to	Jata I
A; Descriptio	on: Production of a Tobacco mosaic virus (TMV) inactivating neotop specific
A; Reference	5057
A; Accession:	S25058
A;Status: pr	reliminary
A; Molecule t	ype: mRNA
A; Residues:	I-235 <fis></fis>
A; Cross-refe	rences: EMBL:X67211; NID:g54828; PIDN:CAA47650.1; PID:g54829
C;Superfamil	y: immunoglobulin V region; immunoglobulin homology
F;38-111/Domain: immune	nain: immunoglobulin homology <imm></imm>
•	
Query Match	75.9%;
Best Local	Similarity 79.3%; Pred. No. 5.2e-33;
Matches	a)
ò	2 IELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPARL 61
Dp 5	4 IVLTQSPAIMSA
3	62 - SECRETARIA TARANTA TANANTA TARANTA TARANT
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84 SGSGSGTSYSLTISSMEAEDAATYYCQQWSSNPLTFGAGTKLELKRADAAPTVSIF 139

anti-phosphorylch 21-Jan-2000 RESULT 2
PLO013
Ig kappa chain precursor V region (4C11) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus-1992 #sequence_revision 30-Jun-1992 #text_change 21-Jan-;
C;Accession: PLO013
R;Cheng, H.L.; Sood, A.K.; Ward, R.E.; Kieber-Emmons, T.; Kohler, H.
Mol. Immunol. 25, 33-40, 1988
A;Title: Structural basis of stimulatory anti-idiotypic antibodies.
A;Reference number: PL0011; MUID:88142863; PMID:3125424
A;Reference number: PL0011; MUID:88142863; PMID:3125424
A;Residues: 1-140 cHE>
A;Experimental source: cell line 4C11
C;Comment: This protein is an anti-idiotypic antibody that induces an a:
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;1-22/Domain: signal sequence #status predicted cSIG>
F;23-129/Product: Ig heavy chain V region (4C11) #status predicted cMAT
F;46-55/Region: complementarity-determining 1 an

<MAT>

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A30562
Ig kappa chain V
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C; Date: 31-Dec-1991 #sequence revision 09-Aug-1996 #text_change C; Accession: JL0079; A49044; E49044
R; Kaartinen, M.; Rocca-Serra, J.; Maekelae, O.
R; Kaartinen, M.; Rocca-Serra, J.; Maekelae, O.
Mol. Immunol. 25, 859-865, 1988
A; Title: Combinatorial association of V genes: one VH gene codes A; Reference number: JL0076; MuID: 89096973; PMID:3211160
A; Rccession: JL0079
A; Roccession: JL0079
A; Note: the authors translated the codon TTC for residue 8 as Pr A; Note: the codon given for 1-Met (AGT) is inconsistent with the R; Milstein, C.; Even, J.; Jarvis, J.M.; Gonzalez-Fernandez, A.; Eur. J. Immunol. 22, 1627-1634, 1992
A; Rilstein, C.; Even, J.; Jarvis, J.M.; Gonzalez-Fernandez, A.; Eur. J. Immunol. 22, 1627-1634, 1992
A; Rilstein, C.; Even, J.; Jarvis, J.M.; Gonzalez-Fernandez, A.; Reference number: A49044
A; Molecule type: DNA
A; Residues: 1-25 < MIL>
A; Roccession: A49044
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C; Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-89/Domain: immunoglobulin homology <IMM>
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                                                                          Score 448; DB 2; I
Pred. No. 3.5e-32;
3; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 437; DB 2;
Pred. No. 2.5e-31;
9; Mismatches 14
                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V region (F3, anti-AFP) - mouse (F: C; Species: Mus musculus (house mouse)
C; Date: 10-Nov-1997 #sequence_revision 23-Jan-1991
C; Accession: PC4405
R; Deng, J.B.; Han, H.; Su, C.Z.; Chen, C.Q.
Chinese Biochem. J. 12, 648-653, 1996
A; Title: Generation of a phage display library of A; Reference number: PC4405
A; Reference number: PC4405
A; Molecule type: mRNA
A; Residues: 1-107 < DEN>
A; Cross-references: UNIPROT: Q8K1F0
;71-77/Region: complementarity-determining 2;110-118/Region: complementarity-determining;130-140/Domain: constant region (fragment)
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ilarity 78.5%;
Conservative
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Best Local Similarity 75.9%;
Matches 88; Conservative
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Best Local Similarity
Matches 84; Conser
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immunofluore
A;Note: sequence extracted from NCBI backbone (NCBIN:106802, NCBIP:106809)
A;Accession: B49044
A;Accession: B49044
A;Molecule type: DNA
A;Residues: 114-116 km1L2>
A;Cross-references: GB:837664; NID:9250215; PIDN:AAB22332.1; PID:9250218
A;Experimental source: BALB/c germ-line
A;Note: sequences extracted from NCBI backbone; partial nucleotide sequences of 13 ad
A;Note: sequence extracted from NCBI backbone (NCBIN:106807, NCBIP:106822)
C;Genetics:
A;Gene: V(kappa)Ox1
A;Introns: 17/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (C;Complex: An immunoglobulin heterotetramer; such as IgA and IgM, the subunits associate into C;Superfamily: immunoglobulin immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;1-22/Domain: signal sequence #status predicted <SIG>F;23-130/Product: Ig kappa chain V region (6F6) #status predicted <MAT>F;45-109/Disulfide bonds: #status predicted
F;45-109/Disulfide bonds: #status predicted
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C;Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_change 21-Jan-2000
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C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 21-C;Accession: B54378
R;Agazie, Y.M.; Lee, J.S.; Burkholder, G.D.
J. Biol. Chem. 269, 7019-7023, 1994
A;Title: Characterization of a new monoclonal antibody to triplex D A;Reference number: A54378; MUID:94165109; PMID:7509814
A;Accession: B54378
A;Cross-references: GB:S68985; NID:g545746; PIDN:AAB30096.1; PID:g5A;Cross-references: GB:S68985; NID:g545746; PIDN:AAB30096.1; PID:g5A;Cross-reference extracted from NCBI backbone (NCBIN:144174, NCBIP:C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
F;16-89/Domain: immunoglobulin homology <IMM>
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SGSGSGTSYSLPISRMEAEDAATYYCQQRSSYPITFGAGTKLELK
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Pred. No. 4.9e-30;
3; Mismatches 15;
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2.2e-30;
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Pred. No.
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Date: 06-Oct-1994 #sequence_revision
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34; Conservative
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ilarity 79.6%;
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81; Conservative 1
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-88/Domain: immunoglobulin homology <IMM>
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                   P.N.; Bhattacha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Mus musculus (house mouse)
C;Date: 10-Jul-1998 #sequence_revision 10-Jul-1998 #text_change
C;Accession: PD0011
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10; Mismatches 11
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                    F.; Akolkar,
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Chinese Biochem. J. 12, 648-653, 1996
A;Title: Generation of a phage display library
A;Reference number: PC4405
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C; Accession: A30562
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Ako. J. Immunol. 142, 888-893, 1989
A; Title: Amino acid substitutions in V-H CD. A; Reference number: A30562; MUID:89110066; A; Accession: A30562
A; Accession: A30562
A; Molecule type: mRNA
A; Molecule type: mRNA
C; Superfamily: immunoglobulin V region; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
F; 16-89/Domain: immunoglobulin homology <IM
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C;Date: 06-Jan-1995 #sequence_revision 06-
C;Accession: S29591
R;Kavaler, J.
submitted to the EMBL Data Library, April
A;Reference number: S26459
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A;Residues: 1-107 <DEN>
C;Superfamily: immunoglobulin V region;
F;16-89/Domain: immunoglobulin homology
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A;Accession: S29591
A;Status: preliminary
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C; Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 21-Ja
C; Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 21-Ja
C; Accession: PS0071
R; Levy, S.; Campbell, M.J.; Levy, R.
J. Exp. Med. 170, 1-13, 1989
A; Title: Functional immunoglobulin light chain genes are replaced by A; Reference number: A92781; MUID:89310348; PMID:2501443
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-89/Domain: immunoglobulin homology <IMM>
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Pred. No. 1.1e-29;
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immunoglobulin homology
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S11119
Ig kappa chain V region (clone NQ6-8.3.1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_chang
C;Accession: S11119
R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A;Title: mRNA sequences define an unusually restricted IgG res
A;Reference number: S07331; MUID:83271467; PMID:6877353
A;Residues: 1-107 <NAT>
C;Superfamily: immunoglobulin V region; immunoglobulin homolog
F;16-89/Domain: immunoglobulin homology <IMM>
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                                                                                                               Ig kappa chain V region (anti-idiotypic) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_cha
C; Accession: B49049
R; Armandola, E.A.; Mariani, S.M.; Zwickl, M.; Hardman, N.; F
Eur. J. Immunol. 22, 2893-2899, 1992
A; Title: Molecular analysis of anti-idiotypic monoclonal ant
A; Reference number: A49049; MUID: 93049629; PMID: 1425914
A; Accession: B49049
A; Reference number: B49049
A; Residues: 1-104 < ARM>
A; Residues: 1-104 < ARM>
A; Residues: 1-104 < ARM>
C; Superfamily: immunoglobulin V region; immunoglobulin homolog; Superfamily: immunoglobulin homolog; Keywords: immunoglobulin homolog; Keywords: immunoglobulin homolog; Keywords: immunoglobulin homology < IMM>
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                    mouse (fragment)
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red. No. 2e-29;
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S11118
Ig kappa chain V region (clone NQ5-61.1.2) - mouse C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 18-Jul-1997
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Pred. No. 3e-29;
; Mismatches
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Similarity 76.4%;
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C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 09-09.
C; Date: 15-Dec-1988 #sequence_revision 15-Dec-1988 #text_change 09-09.
C; Accession: G27887
R; Caton, A.J.; Brownlee, G.G.; Staudt, L.M.; Gerhard, W.
EMBO J. 5, 1577-1587, 1986
A; Title: Structural and functional implications of a restricted ant A; Reference number: A91043; MUID:86300658; PMID:2427335
A; Reference number: A91043; MUID:86300658; PMID:2427335
A; Reference number: A91043; MUID:86300658; PMID:2427335
A; Residues: 1-106 < CAT>
C; Comment: This sequence was determined from the germline gene C; Comment: This chain was isolated from a hybridoma protein that bic C; Superfamily: immunoglobulin V region; immunoglobulin c; Keywords: heterotetramer; immunoglobulin
F; 16-89/Domain: immunoglobulin homology < IMM>
F; 16-89/Domain: immunoglobulin homology < IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_ch
C; Date: 23-Mar-1989 #sequence_revision 23-Mar-1989 #text_ch
C; Accession: B30562
R; Sikder, S.K.; Borden, P.; Gruezo, F.; Akolkar, P.N.; Bhat
J. Immunol. 142, 888-893, 1989
A; Title: Amino acid substitutions in V-H CDR2 change the ic
A; Reference number: A30562; MUID:89110066; PMID:2464031
A; Accession: B30562
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-107 <SIK>
C; Superfamily: immunoglobulin V region; immunoglobulin homc
C; Keywords: heterotetramer; immunoglobulin
F; 16-89/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                revision 23-Mar-1989 #text
                                  15;
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Pred. No. 1.4e-29;
); Mismatches 17;
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Pred. No. 1.6e-29;
7. Mismatches 18;
ced. No. 1.3e-29;
Mismatches 15
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Search completed: October 13, 2004, 13:00:06 Job time: 26.2958 secs

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Compugen Ltd
GenCore version (c) 1993 - 2004
         Copyright
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- protein search, using sw model OM protein

Sec 2004, 12:47:33 October 13, Run on:

; Search time 119 Se. (without alignments) 575.375 Million cell

updates/sec

US-10-049-868A-3

score: Title: Perfect

.....GTKLELKRADAAPTVS 606 1 DIELTQSPAIMSASPGEKVT... Sequence:

IFKL

BLOSUM62 Gapop 10.0 , Scoring table:

1825181 seqs, 575374646 residues Searched:

0.5

Gapext

1825181 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 200000000 DB DB Minimum Maximum

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_tremb1:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARTES

		, 0 4			SUMMARIES	
Result		Query				
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30	•		3	0	<u> Q811</u> <u>C</u> 3	1c3 mus mus
31	S	60.4	97	7		ar11055 mus

Q91W12 PRELIMINARY; PRT; 235 AA.
Q91W12;
Q91W12;
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus.
NCBI_TaxID=10090;

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RESULT Q91W12

Aar11066 mus muscu	Aar11062 mus muscu	r10983 mus	nm srum	ราม ธาน	mus mu		anu snw		0.2	2 mus	fpih7 homo sapi	9 homo sapi	ah34141 homo sap
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C QRKIF2;

T 01-OCT-2002 (TrEMBLrel. 22, Created)

T 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)

T 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)

T 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)

T 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)

Anti-VIPase light chain variable region (Fragment).

Mus musculus (Mouse).

E barayota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

NCBI TaxID=10090;

NCBI TaxID=10090;
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Pred. No. 5.5e-38;
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Q8K1F0;
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                              TISSUE=Mammary tumor. WAP-TGF alpha model. 7 months old;

MEDLINE=22388257; PubMed=12477932;

A REDLINE=22388257; PubMed=12477932;

A Strausberg R.D., Feingold E.A., Grouse L.H., Derge J.G.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko D., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Mitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

A Jones S.J., Marra M.A.;

A Jones S.J., Marra M.A.;
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Ξ.,
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Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Muri
NCBI_TaxID=10090;
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Last sequence update)
Last annotation update)
chain (Fragment).
                                                                                                                                                                                                                                                                                 sequences.";
id. Sci. U.S.A. 99:16899-16903(2002)
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Pred. No. 1.6e-37,
9; Mismatches 18
                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=mix FVB/N;
TISSUE=Mammary tumor. WAP-TGF alpha model. 7
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6
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Submitted (APR-2001) to the EM EMBL; BC006643; AAH06643.1; -.. PDB; 1AY1; X-ray; L=23-232.
PDB; 1HQ4; X-ray; A/C=23-235.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_WHC.
InterPro; IPR003596; Ig_W.
Pfam; PF07654; C1-set; 1.
Pfam; PF00047; ig; 1.
SMART; SM00406; IG_NHC; UNKN
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Similarity 76.7%;
89; Conservative 9
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(TrEMBLrel. 20, 1
(TrEMBLrel. 26, 1
2 variable light 0
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Proc. Natl. Acad.
                 FROM N.A.
    [1]
SEQUENCE FROM N.A
STRAIN=mix FVB/N;
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Q8VDD0;
01-MAR-2002 (T
01-MAR-2002 (T
01-MAR-2004 (T
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Best Local S
Matches 89
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SEQUENCE FROM N.A.

STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;

Zhou Y.-X., Taguchi H., Planque S., Karle S., Nishiyama Y.,

Zhou Y.-X., Taguchi H., Planque S., Karle S., Nishiyama Y.,

EMBL; AF516285; AAM64203.1; -.

R PIR; A33933; A33933.

R PIR; PC4405; PC4405.

R PIR; PC4405; PC4405.

R PIR; PC4405; PC1406.

IN InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

INTERPRO; IPR003596; Ig_v.

SRART; SM00406; IGv; 1.

SRART; SM00406; IGv; 1.
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17; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6;
18;
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Pred, No. 4.7e-36
                                                                                                                                                                                                                                                                                                                          Score 444; DB 2;
Pred. No. 4.4e-37;
); Mismatches 17
                                                                                                                              EMBL/GenBank/DDBJ
                                              the EMBL/GenBank/DDBJ
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Mismatches
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11901 MW;
                                                           SEQUENCE FROM N.A.
STRAIN=BALB/c;
Chernajovsky Y.;
Submitted (OCT-2001) to the EMEMBL; AJ41631; CAC94866.1; -.
PIR; G27887; G27887.
HSSP; P01834; 1MIM.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
NON_TER
                                                                                                                                                                                                                                                                                            14525 MW;
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77.1%;
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Similarity 76.6%;
85; Conservative
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                                                40
                                Sembi P.;
Submitted (JAN-2002)
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112 AA;
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134 AA;
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84; Conser
SEQUENCE FROM N.A STRAIN=BALB/c;
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59 ARLSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAP
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SMART; SM00406; iGv
PROSITE; PS50835; I
  Eukaryota; Metazoa
Mammalia; Eutheria
NCBI_TaxID=10090;
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les 83; Conser
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114
114 A
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Q8K1F1;
01-OCT-2002
01-OCT-2002
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Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG response phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
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SGSGNSYSLTISSTEGEDAATYYCQQWSSNPPTFGGGTKLEIKRADAAP
                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-ror send an email to license@isb-sib.ch).
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Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
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Framework-4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6F694284ECFA68E6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 421; DB 1;
Pred. No. 7.4e-35;
9; Mismatches 15
                                                                13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-VI region NQ2-17.4.1.
Mus musculus (Mouse).
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                                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11561 MW;
                                                                                                                                                                                                                                                                                                                        EMBL; K00735; AAA38680.1; -. HSSP; P01679; 2FBJ. InterPro; IPR007110; Ig-like InterPro; IPR003596; Ig-v.
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LIKE;
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                                                 STANDARD;
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SMART; SM00406; IGv;
PROSITE; PS50835; IG
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82; Conser
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01-OCT-2002
01-OCT-2002
                                                 MOUSE
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NON_TER
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                                                 KV6F
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID=10090;
Euteleostomi
Murinae; Mu
                                                                                                             Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S., Nishiyama Y., Pau Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF516284; AAM64202.1; -.
RPIR; A33933; A33933.
RPIR; PH1058; PH1058.
R PIR; PH1058; PH1058.
R HSSP; P01837; 25C8.
R InterPro; IPR007110; Ig-like.
R InterPro; IPR003596; Ig-v.
R InterPro; IPR003596; Ig-v.
R Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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databases.
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Craniata; Vertebrata; E
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-VIPase light chain variable region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                SEQUENCE FROM N.A.

STRAIN=BALB/c; TISSUE=Hyperimmunized spleen;
Zhou Y.-X., Taguchi H., Planque S., Karle S., Ni:
Submitted (MAY-2002) to the EMBL/GenBank/DDBJ date EMBL; AF516282; AAM64200.1; --
PIR; A33933; A33933.
PIR; PH1085; PH1085.
HSSP; P01837; 25C8.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                     Score 419; DB 2;
Pred. No. 1.2e-34
2; Mismatches 18
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Pred. No. 1.3e-34;
Mismatches 20
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Chordata;
Rodentia;
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12017 MW;
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Similarity 72.3%;
81; Conservative 12
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ilarity 72.8%;
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IG_LIKE;
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(Wonse)
Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                          BI_TaxID=10090;
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01-OCT-2003 (
01-OCT-2003 (
01-MAR-2004 (
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SEQUENCE
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          114
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                                                                                           Ig kappa chain V-VI region NQ6-8.3.1.

Mus musculus (Mouse).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

NCBI_TaxID=10090;
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ARFSGSGSGTSYSLTISSMEAEDAATYYCQQYHSYPRTFGGGTKLEIKE
                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG respons phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Ithe European Bioinformatics Institute. There are no restuse by non-profit institutions as long as its content modified and this statement is not removed. Usage by are entities requires a license agreement (See http://www.isbor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Framework-1.
Complementarity-determining-Framework-2.
Complementarity-determining-Framework-3.
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Complementarity-determining.
Framework-4.
By similarity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           CRC64;
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Pred. No. 2.4e-34;
); Mismatches 16
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on update)
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(Rel. 05, Last sequence update)
(Rel. 38, Last annotation update)
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-AUG-1987 (Rel. 05, Last sequence up
-JUL-1999 (Rel. 38, Last annotation kappa chain V-VI region NQ5-61.1.2.
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ilarity 76.4%;
Conservative
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InterPro; IPR003596; Ig_v.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE;
Hybridoma; Immunoglobulin
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HSSP; P01679; 2FBJ.
                                                     STANDARD;
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13-AUG-1987
13-AUG-1987
15-JUL-1999
IG kappa chai
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13-AUG-1987
15-JUL-1999
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KV6H_MOUSE
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                                                     MOUSE
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Best Local S
Matches 81
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KV6I
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Sciurognathi; Muridae; Musinae; Mus
  Murinae; Mus
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 Euteleostomi
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Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity-
                                                                     SEQUENCE FROM N.A.
MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG response phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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Framework-1.
Complementarity-determining-1
Framework-2.
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TISSUE=Mammary tumor metastatized to lung. MMTV-LTR/Wn
Expression driven by an MMTV-LTR enhancer.;
MEDLINE=22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CRC64
Craniata; Vertebrata; E
Sciurognathi; Muridae;
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Last sequence update)
Last annotation update)
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Pred. No. 3e-34;
; Mismatches
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Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                          EMBL; K00739; AAA38684.1; -. HSSP; P01679; 2FBJ.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Hybridoma; Immunoglobulin V re-
Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11605 MW;
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81; Conservative
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Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Raha S.S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Jones S.J., Marra M.A.,
T. "Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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MEDLINE=83271467; PubMed=6877353;
Kaartinen M., Griffiths G.M., Markham A.F., Milstein C.;
"mRNA sequences define an unusually restricted IgG response phenyloxazolone and its early diversification.";
Nature 304:320-324(1983).
-!- FUNCTION: Anti-2-phenyl oxazolone (PHOX) Antibody.
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                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN=CZECH II;
TISSUE=Mammary tumor metastatized to lung. MMTV-LTR/Wnt1
Expression driven by an MMTV-LTR enhancer.;
Strausberg R.;
Strausberg R.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                   sequences.";
ad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 415; DB 2;
Pred. No. 7.4e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P04941;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                           Submitted (AUG-2003) to the EMBL/GenBarber, BC055911; AAH55911.1; -... InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig c1.
InterPro; IPR003006; Ig MHC.
InterPro; IPR003596; Ig v.
Pfam; PF07654; C1-set; I.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS00290; IG LIKE; 2.
PROSITE; PS00290; IG MHC; UNKNOWN I.
Hypothetical protein.
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72.8%;
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235 AA; 2
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Mus musculus (Mouse)
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Proc. Natl. Aca
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between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Framework-1.
Complementarity-determining-1
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Complementarity-determining-
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Complementarity-determining
Framework-4.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
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No. 3.8e-34
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Pred. No. 7.9e-34
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Hybridoma; Immunoglobulin V re
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76.4%;
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71.8%;
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nae; Mus.
                                                                                                              Mus musculus (Mouse). --
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID=10090;
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02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murin
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                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.,
"Antinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY436822; AAR10982.1; -.
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STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.
"Antinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases
EMBL; AY436834; AAR10994.1; -.
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11819 MW; EE2F666368BC91BC CRC64;
                        AAR10982
AAR10982;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
ANA immunoglobulin kappa light chain (Fragment).
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Pred. No. 7.9e-34;
5; Mismatches 18;
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79; Conserv
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77; Conser
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Best Local S
Matches 79
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mu
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                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN=B6.Sle1; TISSUE=Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Zhou J.X.,
"Antinuclear autoantibodies from B6.Sle1 mice.";
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases
EMBL; AY436888; AAR11048.1; -.
NON TER
108
108
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RESULT 15
AAR11048
ID AAR11048
AC AAR11048;
C AAR11048;
DT 02-MAR-2004 (TrEMBLrel. 27, Created)
DT 02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
DT 02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
DS ANA immunoglobulin kappa light chain (Fragment).
OS Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 410; DB 2; I
Pred. No. 9.7e-34;
3; Mismatches 17;
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Job time : 120 secs
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[larity 72.0%;
Conservative 13
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108 AA;
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Best Local S
Matches 77
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5.1.6
Compugen Ltd.
version
- 2004
GenCore (c) 1993
       Copyright
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protein search, using sw model OM protein

Seconds 2004, 12:44:23 October 13, on: Run

updates/sec ; Search time 111.06 (without alignments) 384.353 Million cell

..GTKLELKRADAAPTVS DIELTQSPAIMSASPGEKVT.. 909 score: Sequence: Title: Perfect

US-10-049-868A-3

119

IFKL

BLOSUM62 Gapop 10.0 Scoring table:

0.5 Gapext

Searched:

2002273 of hits satisfying chosen parameters: Total number

2002273 seqs, 358729299 residues

2000000000 length: length: sed DB DB Minimum Maximum

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

A_Geneseq_23Sep04:*
geneseqp1990s:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Query Result No.

cription	4 1018	m79827 Mouse	r08347 Ligh	b75616 Anti-	w41392 Chime	ar21301 Murine	do43520 Amino	r43674 Mouse a	r99644 Anti-h	97377 Murine	r21293 Murine	ar13060 Monocle	br62010 Single	abiliaga m	ab74199 PelR-c	ab70769 Expressi	7000 E Cayot	avises d: card	ae37758 Murine T	T OCITION TO STATE OF THE STATE	av32086 cindle	TOTACA MITTER V	SECTION FILE TILE	dh77307 Veset	object of the contract	aD / 064
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ALIGNMENTS

215 protein; ADE06764 standard; RESULT 1 ADE06764

AA

entry) (first 29-JAN-2004 ADE06764;

light

ID NO:34

protein SEQ

chain

hybrid polypeptide; protein aggregation; prion polypeptide; neuroprotective; nootropic; antidiabetic; anticonvulsant; cerebroprotective; antiparkinsonian; cytostatic; nephrotropic; cardiant; antiinflammatory; antiarteriosclerotic; gene therapy; Creutzfeldt-Jakob disease; scrapie and bovine spongiform encephalopathy; Alzheimer's disease; Type II diabetes; Huntington's disease; amyotrophic lateral sclerosis; amyloidosis; chronic inflammatory disease; erontotemporal dementia; multiple myeloma; plasma cell dyscrasia; familial amyloidotic polyneuropathy; medullary carcinoma; chronic renal failure; congestive heart failure; chronic inflammation;

Synthetic

atherosclerosis

WO2003085086-A2

16-0CT-2003

2003WO-US010856. 08-APR-2003;

2002US-0371610P 09-APR-2002;

INST SCRIPPS RES ^ (SCRI

Moroncini G; Williamson RA, Burton DR,

hybrid polypeptides binding to the infectious form of a diagnosing or treating diseases of protein aggregation s.g. amyloidosis, Alzheimer's disease, renal failure or WPI; 2003-877028/81 N-PSDB; ADE06763. New motif-grafted prion, useful for or conformation, e

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English 115pp; 34; SEQ ID NO Disclosure;

e.g

diabetes.

(a) a hybrid polypeptide (I) comprising: invention describes The present

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2003WO-JP013063

10-OCT-2003;

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a polypeptide motif containing a sufficient number of contiguous amino acid residues from a polypeptide associated with a disease of protein pelypeptide or to a disease-associate conformer of the polypeptide; and polypeptide or to a disease-associate conformer of the polypeptide; and polypeptide from which the motif is derived, where the resulting hybrid polypeptide binds with greater affinity to a disease causing or infectious conformer of the polypeptide that is the resulting hybrid polypeptide motif compared to a benign form of the polypeptide. Also escribed: (1) a mucleic acid molecule, (3) a cell comprising the vector; (4) detecting an isoform or a PrPSc form of a prion polypeptide or a comprising an isoform or a PrPSc form of a prion polypeptide or a propertied sassociated with a disease of protein aggregation, in a sample; (5) a solid support comprising a plurality of polypeptides described above; (6) detecting cells that contain a protein conformer associated with a disease of protein aggregation, (7) preparing a hybrid molecule that specifically interacts with one conformer of a protein involved in the disease mentioned above; and (8) an anti-idiotype antibody that specifically binds to an infectious form of a protein involved in the disease mentioned above; and (8) an anti-idiotype antibody that specifically binds to an infectious form of a protein conformer of a protein. (1) has neuroprotective, antiparkinsonian, cytostatic, nephrotropic, cardiant, cerebroprotective, antiparkinsonian, cytostatic, nephrotropic, cardiant, antiparkinsonian, cytostatic, nephrotropic, antiparkinsonian, conformation, such as Creutzfelde-Jakob disease, scrapic and bovine spongilorm encephalopsthy, Alzheimer's disease, reactive amyloidosis, reactive amyloidosis, reactive amyloidosis, sesociated with chronic inflammatory disease, perceptions and perception of perception and methods a disease, perception and methods of the present invertion and methods a disease, perception of a disease, immunoglobulish amyloidosis, reactive a
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medullary carcinoma
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215 AA; Sequence

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cancer;
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                                           1 DIELTOSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPAR
                                                          Gaps
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G2b light
                                                                                                                                                                                                                                                            antibody;
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                                                                                                                                                                                                                                                                                   ease:
 215;
                                                                                                                                                                                                                                                           cell death-inducing agent; human leukocyte antigen; HLA; 2D7 antibody degradation product; cell proliferation inhibition; antitumour; autoimmune discytostatic; immunosuppressive; antiinflammatory; dermatol Crohn's disease; systemic lupus erythematosus; mouse; IgG
                                                                                        LSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRA
                       Indels
 Length
Score 471; DB 7; I
Pred. No. 2.7e-29;
); Mismatches 17;
                                                                                                                                                                                                                                        Mouse IgG2b light chain protein SEQ ID NO:4.
                                                                                                                                                                       ADM79827 standard; protein; 144
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77.7%;
ilarity 77.8%;
Conservative
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              Similarity
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                        91;
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entry)

(revised) (first ent

25-MAR-2003 05-MAR-1991

WO2004033499-A1

22-APR-2004.

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The present invention describes cell death-inducing agents containing the degradation product of an antibody which can recognise human leukocyte antigen (HLA). Also described: (1) a degradation product of the 2D7 antibody; (2) a similar degradation product of an antibody which is: (a) cone containing an amino acid sequence of (SEQ ID NO:6) but with some amino acids substituted, deleted, inserted and/or added and being functionally equivalent to the parent; (c) one containing the CDR amino-acid sequences of (SEQ ID NO:2) and (SEQ ID NO:4) with 18 and 144 amino-acids sequences of (SEQ ID NO:4) but with some amino acids substituted, deleted, inserted and/or added and being functionally equivalent to the parent; (3) a process for producing an antibody with an increase of activity by degrading a HLA-recognising antibody, or by degrading 2D7 antibody; (4) cell death-inducing agents containing the degradation products of antibody, comprising the produced antibodies or 2D7 antibody as active ingredient; (5) agents for inhibiting cell proliferation containing the degradation products of antibody, comprising the produced antibodies or 2D7 antibody; and (7) remedies for autoimmune diseases containing the degradation products of antibody, comprising the produced antibodies or 2D7 antibody; and (7) remedies for autoimmune diseases containing the degradation products of antibody, comprising the produced antibodies or 2D7 antibody; and (7) remedies for autoimmune diseases for treating e-grentci immunosuppressive, antiinflammatory and dermatological activities. The cell death-inducing agents are applicable in antitumour agents and streams and systemic lupus erythematosus. The present sequence containing which is and systemic lupus erythematosus. The present sequence
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Crohn's disease and systemic lupus erythematosus. Th
represents a mouse IgG2b light chain, which is used
exemplification of the present invention.
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17;
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Pred. No.
                                                                                                                                                            Kimura
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77.68;
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                                                                                                                                                                                                                                                                                                                                                 The MAb recognises the epitope on alpha/beta TCR and includes a mediator resonse. It is highly effective in clinical application before, after and during bone marrow transplant surgery, in treatment of cancer and autoimmune diseases, and has applications in immunoregulation. Abs may also be used in detection of immunocompetent T-cells. (Updated on 25-MAR-2003 to correct PA field.)
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                                                                                                                                                                                                                                                                                                antibodies - used for as therapeutic agent
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                           cancer; autoimmune
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                                                                                                                                                                                                                                                                                             Anti-human alpha-beta T-cell receptor monoclonal immunosuppressive therapy in transplantation and immuno-regulation, e.g. to treat cancer.
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         the BMA-031 MAb
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1.3e-28;
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Pred. No. 1.3e-
8; Mismatches
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N-PSDB; AAQ06956, AAQ06958.
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                         alpha/beta T-cell
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BEHRINGWERKE AG
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Matches 90
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The present sequence is that of the light chain variable region of claimed anti-human RANK ligand (RANK-L) murine monoclonal antibody (mAb) 19H22 (isotype IgG2b/Kappa). The mAb is produced by hybridoma cells obtained by immunising CB6 fl mice with human RANK-L protein, and fusing obtained by munising CB6 fl mice with human RANK-L protein, and fusing speen cells with X63-Ag8.653 myeloma cells. 19H22 mAb is specific for human RANK-L, having a binding affinity characterised by a dissociation constant of about 10 power -10 M. RANK-L is a member of the tumour necrosis factor family of proteins and a key regulator of the immune system, bone development and homeostasis. Including Fab or F(ab/)2 chimeric and humanised 19H22 antibodies, including Fab or F(ab/)2 fragments, antibodies comprising the light and heavy chain variable regions of 19H22, and antibodies comprising the complementarity determining regions of 19H22. These are used in methods for the determining regions of 19H22. These are used in methods for the including psoriasis, insulin dependent diabetes, inflammatory bowel including psoriasis, insulin dependent diabetes, inflammatory bowel recombinant production of the antibodies, using isolated nucleic acids and transfected host (preferably mammalian) host cells, and a method of diagnosing conditions associated with Th1 T-cell activity or osteoclast development and activation, in particular those listed above
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complementarity determining region; CDK; INEUMALOIA ALTERITYS; osteoarthritis; osteoporosis; bone cancer; metastasis; osteolysis; osteoarthritis; psoriasis; inflammatory bowel disease; multiple sclerosis; diabetes; cytostatic; antiarthritic; antiinflammatory; osteopathic; antidiabetic; antirheumatic; immunosuppressive; antipsoriatic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treating or preventing osteopenic and osteporosis.
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                                                                                                                                                                                                                                          region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                     "complementarity determining
                                                                                                                                                                                                                                                                                   "complementarity determining
                                                                                                                                                                                                                                       determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8;
16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75.7%; Score 459; DB 5; Similarity 76.7%; Pred. No. 1.4e-28; 9; Conservative 11; Mismatches 16
                                                                                                                                                                                                     49. .55
/label= CDR2
/note= "complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              monoclonal antibody 19H22 for tincluding rheumatoid arthritis
                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORP.
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                                                                                                                                                                          CDR1
                                                                                                                                                                                                                                                                    CDR3
                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0226524P
2000US-0230639P
                                                                                                                                                                                                                                                                                                                                                                            2001WO-US026161
                                                                                                                                                                                                                                                                                                                                                                                                                                                       BEECHAM
                                                                                                                                                           24. .33
/label=
                                                                                                                                                                                                                                                  88. .96
/label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tornetta MA,
                                                cytostalle,
antirheumatic; immunose
antirheumatic; immunose
                                                                                                                                                                                          /note=
                                                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             45pp;
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N-PSDB; ABL53530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SMITHKLINE
SMITHKLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7;
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                                                                                                                                                                                                                                                                                                                 WO200215846-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page
                                                                                                               Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                       21-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                           21-AUG-2001;
                                                                                                                                                                                                                                                                                                                                             28-FEB-2002,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89;
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Best Local S
Matches 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SMIK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease
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                                                                                                                                                           Region
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                                                                                                                                                                                                                                                     Region
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                                                                                                                                            Key
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61

IELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPARL

N

8

ligand; RANK-L; monoclonal antibody; mAb; 19H22; antibody;

chain variable region.

Anti-RANK ligand mAb 19H22 light

RANK

coat; pilus;

group IV/VI chain p specific for phOx.

entry)

(first

d

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d

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Fd; bacteriophage; gene III; filamentous; phagemid; capsid; g3p; binding; adsorption; gene VIII; diverse repertoire; specific binding pairs; replicable genetic display package.
                                                 kappa
                                                                                                                                                               Binding-site
                                                                                                                                                                                   Binding-site
                                                                                                                                           Binding-site
                                                                                                                                                                                                                                                                  10-JUL-1990;
                                                                                                                                                                                                                                              23-JAN-1992
                                                                                                                                                                                                                         WO9201047-A
                                                                                                                                                                                                                                                                                                                                                                               Mccafferty
Jackson RH,
                             21-MAY-1992
                                                 Z
                                                                                                              Synthetic
          AAR21301;
                                                                                                                                                                                                                                                                                                                                                 (CAMB-)
(MEDI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     package.
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                                                 Murine
                                                                                                                                 Кеу
ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence is the light chain variable region of the antibody of the invention. The antibody is an anti-CEA (carcinoembryonic antigen) antibody (806.077 Ab). Host cells or transgenic organisms transformed with DNA encoding the antibody, are used to make the antibody or conjugate. The conjugate is used in a medicament suitable for intravenous administration. The conjugate can be used for cancer therapy, selectively killing tumour cells. The antibody can be used for in vivo or in vitro diagnosis of cancer. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  82
  61
                                                                                                                                                                                    therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LSGSGSGTEFTLEISRVKAEDVGVYYCOOLVEYPLTFGAGTKLELKRADAAPTVSIF 117
                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23 DIELTOSPAIMSASPGEKVTITCSASSSVTYMHWFOOKPGTSPKLWIYSTSNLASGVPAR
  SNLASGVPVRF
                                        117
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                                                                                                                                                                                     cancer
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                                                                                                                                                                                    antibody; carcinoembryonic antigen; 806.077 Ab; car
agnosis; complementarity determining region; light
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 IVLTQSPAIMSASPGEKVTMTCSASSSVSYMYWYQQKPGSSPRLLIYDI
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Pred. No. 3.1e-28;
; Mismatches 15
                                                                                                                                                                  chain
                                                                                                                                                                 antibody 806.077 light
                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                       208pp;
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Θ
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                                                                                                                                                                                                                                                                                                                                                                                                                         antigen
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97GB-00003103
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Best Local Similarity 76.9%;
Matches 90; Conservative
                                                                                             standard; protein;
                                                                                                                                                                                                                                                                                                                                                                         Emery
                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       100-101;
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(first en
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                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy of cancer
                                                                                                                                                                                                                                                                                                                                                                        Edge MD,
                                                                                                                                                                                                                                                                                                                                                                                            1997-558987/51
                                                                                                                                                                 anti-CEA
                                                                                                                                                                                               diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        8; Page
                                                                                                                                                                                                                                                                                                                                                     ZENECA
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14-FEB-1997
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02-JUN-1998
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                                                                                                                                                                  Chimeric
                                                                                                                                                                                                                                        Chimeric
                                          62
                       62
                                                                                             AAW41392
                                                                                                                                                                                       Anti-CEA
  \alpha
                                                                                                                 AAW41392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                 cancer
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                                                                                                                                                                                                                                                                                                                                                     (ZENE
                                                                                                                                                                                                                                                                                                                                                                                                                                    and
                                                                                  AAW41392
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protein; 108 AA

AAR21301 ID AAR21301 standard;

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The VK sequence is one of 23 (AAR21286-308) expressed from a single chain from the VK sequence is one of 23 (AAR21286-308) expressed from a single chain from the Ibrary produces a diverse repertoire of antibody completed from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were separately amplified by CR (AAQ23474-84) and ligated into fdCAT2 (AAQ23463) for expression on FC (AAQ23474-84) and ligated into fdCAT2 (AAQ23463) for expression on the phage surface as fusions with gene III. The resulting library of clones was diverse; 23 hapten binding clones were sequenced revealing 8 vH genes (A-H; AAR21286-92). Most clones were VH-B combinations so a further interachical library was prept. W crossing vH-B with the VK repertoire. The resulting library was screened for hapten binding and 24 clones sequenced. 14 new partners (AAR21293-308) for VH-B were identified. Nearly all the VK genes were "No.11ke"; only f, (from the original library) and h, p, q, and r (from the hierarchical library) were visinal library) and h, p, q, and r (from the hierarchical library) were visinal library) and h, p, q, and r (from the hierarchical library) were visinal library and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) were visinal library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical lib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Griffiths AD;
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                           " D-X-G-X-X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Johnson
, Marks
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90GB-00022845.
90GB-00024503.
91GB-00004744.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  s of specific cells with a
                                                                                                                                          CDR2
                                                                                                                                                                                                             CDR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                            90GB-00015198
                                                                      CDR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24; 209pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAMBRIDGE ANTIBODY MED RES COUNCIL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ,
KP,
                                                                                                                                                                   /88. .96
/label=
/note= "
                                                                                                    49. .55
/label=
88. .96
                                      24. .33
/label=
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Holliger K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-056862/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Producing members recombinant host c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 108 AA;
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ر
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19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
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LSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAP
                                                                                                                                                                                                                                                                                                                                                                     chain poly:peptide for binding chain antigen binding portions
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                  English.
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87US-00092110.
89US-00299617.
                                                                                                                                                                                                                                           90US-00512910
                                                                    AAR43674 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                74.4%;
larity 78.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 78pp;
                                                                                                         (revised)
(first en
                                                                                                                                                                                                                                                                                                    ENZON LABS INC
                                                                                                                                                                                                                                                                                                                                                                                                22;
                                                                                                                                                                                                                                                                                                                       Bird RE,
                                                                                                                                                                                                                                                                                                                                         WPI; 1993-367875/46
N-PSDB; AAQ51535.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 91; Conser
                                                                                                                                     Mouse anti-bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                on 25-MAR-2003 to
PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 215 AA;
                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig
                                                                                                                                                                                                                                                            02-SEP-1986;
02-SEP-1987;
19-JAN-1989;
                                                                                                                                                                                    musculus
                                                                                                                                                                                                                                           25-APR-1990;
                                                                                                                                                                                                      US5260203-A.
                                                                                                        25-MAR-2003
23-MAY-1994
                                                                                                                                                                                                                         09~NOV-1993
                                                                                                                                                                                                                                                                                                                     RC,
                                                                                                                                                       Monoclonal
   61
                                                                                                                                                                 therapy;
                                                                                     AAR43674
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                     Single
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Matches
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                                                  RESULT
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                                      SKLASGVPAR
                                                          SKLASGVPAR
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                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              microbial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                or
                                                                                                                                                                                                                       flavour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present
                                                                                                                                                                                                                                                                                                                                                                                                                          New antibodies and polypeptides that recognize microbial lipases enhance the hydrolytic activity of the enzyme, useful for e.g. biocatalysis or bioconversion reactions, in biosensor technology bioremediation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes antibodies or polypeptides for michlipase, which enhance the hydrolytic activity of the enzyme. The polypeptides, antibodies or immunocomplex are useful for biocathes bioconversion reactions (e.g. hydrolysis, interesterification, esterification, alcoholysis, acidolysis or aminolysis); in bion technology; in bioremediation; in preparing fragrances or flavorhemical modification of alcohols, acids, esters or fats; in contemical and food industries; in preparing detergents; or identifying, purifying or measuring microbial lipases. The pressence represents the variable light chain of monoclonal antised is directed against the Candida rugosa lipase.
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                                    Length 108;
                                                                                    FSGSGSGTSYSLTISSMEAEDVATYYCQQWSSNPLTFGAGTKLEIKRA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody BF1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 156;
                                                                                                                                                                                                                                                                                                                                                                     Barachini
                                                                                                                                                                                                                       fragrance;
                                                                           LSGSGSGTEFTLEISRVKAEDVGVYYCOOLVEYPLTFGAGTKLELKRA
                    Indels
             8;
12;
                                                                                                                                                                                                                     antibody; lipase; biocatalysis; bioconversion; fortergent; light chain; monoclonal antibody BF11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                                                                    HR,
 , DB 2;
                                                                                                                                                                                                   variable light chain of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                74.6%; Score 452; DB 8; 75.2%; Pred. No. 6.1e-28
 Score 452; DB
Pred. No. 4.3e
8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                    Koshkaki
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                                                                                                                                                                                                                                                                                                                                                 SAPIENZA
                                                                                                                                           ADO43520 standard; protein; 156 AA
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                                                                                                                                                                                                                                                                                                                                                 LA
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Cernia E;
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                  8;
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74.6%;
illarity 81.5%;
Conservative
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                                                                                                                                                                                  (first entry)
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N-PSDB; ADO43519.
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88; Conserv
          Similarity
                                                                                                                                                                                                                                                                                                                                                                   Biffoni
                                                                                                                                                                                                                                                                                                                                                                           Soro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             156 AA;
                                                                                                                                                                                                                                                                    WO2004039845-A1
                                                                                                                                                                                                                                                   Mus musculus.
                                                                                                                                                                                                                                                                                                          31-OCT-2002;
                                                                                                                                                                                                                                                                                        13-MAY-2004.
                 88;
                                                                                                                                                                                  29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                          ບັ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 88
                                                                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
Query Match
Best Local
Matches 8
                                                                                                                                                                                                                                                                                                                                                                             Palocci
                                                                                                                                                                                                                                                                                                                                                                   Frati
                                                                                                                        RESULT
ADO4352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
antibody; MAD; affinity; binding; antigen; diagnostics imaging; purification; biosensors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAPTVSIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antigen - comprising lightly linked by peptide linker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 215;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                     chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ed. No. 1e-2
Mismatches
                                                                                                                                                                                                                                                                                                                                 growth hormone MAb light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            451;
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9

Wed

rag

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Fd; bacteriophage; gene II
g3p; binding; adsorption;
                                                                                                                                                                                                                                                                                                                                                             1; Fig
                                                                                                                                                                                                                                         ENZON
                                                                                                                                                                       02-SEP-1986;
02-SEP-1987;
19-JAN-1989;
25-APR-1990;
01-APR-1993;
                                                               chain
                                                                                      Mus musculus
                                                                                                                                                    06-JUN-1995;
                                anti-
                                                                                                          US5518889-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1992
25-MAR-2003
13-NOV-1996
                                                                                                                               21-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR21293;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR21293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                      Antibody
                                                                                                                                                                                                                                                              Bird RE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                         (ENZO-)
                                                                 single
                                Murine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
AAR2129
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gg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
                                                                                                                                                                                                                                                                                                                                                                                                                                               The mature heavy chain (AAR99643) and mature light chain (AAR99644) of the mouse anti-bovine growth hormone monoclonal antibody 3C2 can be utilised in novel single chain binding molecules (AAR99645-48), in which the hypervariable regions from IgG1 3C2 MAb are joined by peptide linkers derived from the Fv regions of an IgA class anti- phosphorylcholine myeloma antibody, MCPC-603. The single chain molecules retain the binding specificity of the light and heavy chains and have the advantages of smaller size, greater stability and reduced cost. They can be used in therapy, diagnostics, imaging, purification and biosensors. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                                                                        uding antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                     regions
nsive than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAPTVSIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'n
                                                                                                                                                                                                                                                                                                                                                                    Immuno:purificn. using single binding chain molecule inclubinding parts of antibody light and heavy chain variable seconnected by a linker - is smaller, stabler and less expensemplete antibodies.
                                                                                                  mO
m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                               engineering; single polypeptide chain binding min; light chain; monoclonal antibody; MAb; cowth hormone; bGH; immunoaffinity purification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 451; DB
Pred. No. 1e-2
6; Mismatches
                                                                            light chain
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                                                                                                                                                                                                                                                                                                                                                                                                                            English
   21
                                                                                                                                                                                                                                                                                                                    Hardman
                                                                                                                                                                                                                              86US-00902971.
87US-00092110.
89US-00299617.
90US-00512910.
                                                                            antibody
                                                                                                                                                                                                           95US-00468992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h
Similarity 78.4%;
91; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; protein;
                                                                                                           chain; light chain; me growth hormone; bGH;
  protein;
                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                             18pp;
                                            (revised)
(first en
                                                                                                                                                                                                                                                                                               LABS INC
                                                                                                                                                                                                                                                                                                                   Ladner RC,
                                                                                                                                                                                                                                                                                                                                         1996-333309/33
)B; AAT13734.
                                                                            monoclonal
   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       215 AA;
                                                                                                                                                                                                                                                                                               ENZON
                                                                                                                                                                                                                                                                                                                                                                                                                              Fìg
                                                                                                                                                                                                                              02-SEP-1986;
02-SEP-1987;
19-JAN-1989;
25-APR-1990;
01-APR-1993;
                                                                                                                                                                                                           06-JUN-1995;
                                                                                                                                                                 US5534621-A
                                                                                                                                                                                      09-JUL-1996
                                            25-MAR-2003
11-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      せ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Mac.
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                                                                            Anti-bGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match
                                                                                                                                                                                                                                                                                                                                                                                                                               Example;
                                                                                                                                                                                                                                                                                                                    Bird RE,
 AAR99644
                                                                                                 Antibody
                                                                                                                                                                                                                                                                                                                                         WPI; 199
N-PSDB;
                                                                                                                                                                                                                                                                                               (ENZO-)
                                                                                                                        bovine
                                                                                                                                             Зp
                                                                                                              heavy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR973
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Portions of the heavy chain (AAR97376) and light chain (AAR97377) of murine Igg1 anti-bovine growth hormone monoclonal antibody 3C2 can be incorporated into novel single polypeptide chain binding molecules (see also AAW02188-90). These are expressed in host cells using DNA constructs (see also AAT36460-62) that include heavy and light chain encoding sequences (AAT29056 and AAT29057) joined by linker moleties. Following expression and refolding, the single chain binding molecules show the binding characteristics of the aggregate of the 2 original heavy and light chains of the variable region of the antibody. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pilus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ÉОЧ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LTOSPAIMSASPGEKVTMTCSASSSV--NYMHWFQQESGTFPKRRIYDTSKLASGVPARL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             replacement
easier to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAPTVSIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   coat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chain;
-· BGH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                filamentous; phagemid; capsid; te VIII; diverse repertoire;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 215;
                                                                                        light chair
h hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 დ
დ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunoassay using single chain antigen binding mol. - at labelled or immobilised antibody, are less immunogenic, engineer, more stable and less expensive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chain h specific for phox.
                                                                                        MAb; li
growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 451; DB 2 Pred. No. 1e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ed. No. 1e-2
Mismatches
                                                                                      nonoclonal antibody; immunoassay; bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          108 AA
                                                                                         monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22; 78pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                         봈
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    III; fi
1, gene
                                                     chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9
                                                                                                                                                                                                                                                                                          86US-00902971.
87US-00092110.
89US-00299617.
90US-00512910.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Hardman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             74.4%;
illarity 78.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
                                                                                                                                                                                                                                                         95US-00468988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine VL kappa group IV/VI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                   entry)
                                                     light
                                                                                         engineering; main antibody;
                                                                                                                                                                                                                                                                                                                                                                                                       HNC
(revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                         Ladner RC,
                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-259060/26
N-PSDB; AAT29057.
                                                     BGH MAD
                                                                                                                                                                                                                                                                                                                                                                                                       LABS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 91; Conser
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, (from the library) were type "h". The
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coupled to chicked
rately amplified by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   values found affinit- ies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inding and 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            307, 309-311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Library of a revealing vk genes (
                                                                                                                                                                                                                                                                                                                                                                                                          fiths AD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vn genes (A-H; AAR21264-71) in a variety of pairings with 7 Vk gene; AAR21286-92). Most clones were VH-B combinations so a further hierarchical library was prepd. by "crossing" VH-B with the Vk gene clones sequenced. 14 new partners (AAR21293-308) for VH-B were the clones sequenced. 14 new partners (AAR21293-308) for VH-B were clones sequenced. 14 new partners (AAR21293-308) for VH-B were clones, 15 very and h, p, q, and r (from the hierarchical library) and h, p, q, and r (from the hierarchical library) very coriginal library) and h, p, q, and r (from the hierarchical library) to Vkox1 type genes. Of the 24 hierarchical clones, 4 were of type "h". C kd of VH-B/Vk-d for phOx-GABA was 10 nM, one of the highest values for this suggests that phage bearing scfv fragments having weak affinition and be selected with antigen, probably due to the avidity of the multipody heads on the phage. The different combinations could alone heads.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ld also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              display
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'n
                                                                                                                                                                                                                                                                                                                                                                                                           Grifi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The VK sequence is one of 23 (AAR21286-308) expressed from a fragments specific for 2-phenyl-5-oxazolone (phox). It was personal from mRNA from mice immunised with phox couples serum albumin. The VH and VL kappa sequences were separately pCR (AAQ23474-84) and ligated into fdCAT2 (AAQ23463) for expthe phage surface as fusions with gene III. The resulting liclones was diverse; 23 hapten binding clones were sequenced VH genes (A-H; AAR21264-71) in a variety of pairings with 7; AAR21286-92). Most clones were VH-B combinations so a further archical library was prepd. by "crossing" VH-B with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Producing members of specific binding pairs - by expression recombinant host cells with a secreting replicable genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        heads on the phage. The different combinations cou on a basis of antigen affinity. See also AAR21260-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 108;
    display package
                                                                                                                                                                                                                                                                                                                                                                                                          Hoogenboom HRJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , DB 2;
8.9e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 448; DB Pred. No. 8.9e-8; Mismatches
   genetic
                                                                                                                                                           D-X-G-X-X motif
                                                                                                                                                                                                                                                                                                                                                             ANTIBODY TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                          KS,
                                                           Location/Qualifiers
 replicable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                        Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                      Marks
                                                                                                                                                                                                                                                                       90GB-00015198.
90GB-00022845.
90GB-00024503.
91GB-00004744.
                                                                                                                  CDR2
                                                                                                                                              CDR3
                                                                                      CDR1
                                                                                                                                                                                                                                               90GB-00015198
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80.6%;
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/label=
49. .55
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  pairs;
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/note=
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specific binding
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19-OCT-1990;
12-NOV-1990;
06-MAR-1991;
15-MAY-1991;
                                                                      Binding-site
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Matches 87
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                                                           Key
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The OK3T light chain sequence was deduced from the cDNA sequence isolated from a library prepared from OK3T producing cells. The library was screened with a probe complementary to a region in the mouse kappa constant region. The OK3T sequence was used in CDR- grafting experiments to prepare humanised antibodies. NOTE: Data found in patent WO9109968 has been used to index this entry. (Updated on 25-MAR-2003 to correct PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          with heavy
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comprising CDR grafted antibody vivo therapy and diagnosis.
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                                                                                                                                           OK3T; light chain; humanised antibodies; CDR-grafting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 445; DB 2; I
Pred. No. 3.2e-27;
); Mismatches 19;
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23. .235
                                                                                                                        antibody OK3T light chain
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1. .22
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90WO-GB002017
                                                        protein;
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ilarity 75.0%;
Conservative
                                                                                             (revised)
(first entry)
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light chains, for use in
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/label=
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P-PSDB; AAR13060.
                                                       AAR13060 standard;
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Best Local Similarity
Matches 87; Conser
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21-DEC-1990;
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03-OCT-1991
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                                                                                                                        Monoclonal
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                                                                                                                                                                                         Peptide
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                                    AAR13060
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SKLASGVPAR

DIELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDT

LSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRA 1

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SKLASGVPAR

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HOFFMANN LA
coli
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 255 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
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                                                                              26-APR-1999;
                                  EP1048732-A1
                                                                                                    26-APR-1999;
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Escherichia
                                                         02-NOV-2000
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Best Local S
Matches 88
           Synthetic
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AAB74199
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                                                                     gene therapy;
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useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA
                                                                                                                                                                                                                                                                                              Increasing the content of transgene-coded biomolecules in animal, useful for producing proteins for diagnosing, prevetreating viral diseases and cancer, comprises changing the of ATP and/or ADP.
                                                                                                                                                                                                                                                                                                                                                                          The invention relates to increasing the content of one or coded biomolecules in an organism and involves changing the of ATP and/or ADP in cells of the organism. The yield of twolecules in host cells is often insufficient for industrium the method increases the yield of transgenic molecules in plant host cells, therefore facilitating their production industrial scale. The proteins produced by the method are diagnosing, preventing and/or treating viral diseases and present sequence represents a single-chain (scFv) antibody exemplify the increase in the expression of scFv antibodie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              coli expression plasmid pUBS520.ScFvOx encoded protein
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                                                                     virucide;
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                                                                     biomolecule; ATP; ADP; cytostatic; nerapy; scFv; antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 444; DB 6 Pred. No. 4e-27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7;
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disulfide bridge.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73.3%;
illarity 80.0%;
Conservative
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                                                                                                                                                                                             2001EP-00130319
                                                                                                                                                                         2001EP-00130319
                                                antibody
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                                                                                                                                                                                                                                          Petsch
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                                                                                antisense therapy; scFv;
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N-PSDB; ACC84876.
                        (first
                                                (scFv)
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                                                                                                                                                                                                                    DUERING K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tubers
                                                                                                                                                                        19-DEC-2001;
                                                                                                                                                                                             19-DEC-2001;
                                               Single-chain
                                                                                                                             EP1321524-A1
                                                                                                                                                 25-JUN-2003.
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                                                                      Transgenic;
                          -OCT-2003
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Best Local S
Matches 88
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                                                                                                      Synthetic
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   ABR62010;
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This invention describes a novel preparation of a water-soluble, naturally occurring eukaryotic polypeptide containing two or more cysteine units bound via a disulfide bridge which comprises cultivation of prokaryotic cells in the presence of arginine or an amide compound. The method is useful for the preparation of eukaryotic proteins e.g. proteases, interferons, protein hormones, antibodies or antibody fragments (e.g. a single chain FV fragment that binds to thyroid stimulating hormone). It is especially useful for preparing proteins with more than five disulfide bridges, e.g. recombinant plasminogen activator (rPA). The technique is simple and does not require in vitro after-treatment, such as the removal of inclusion bodies, reduction or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DIELTOSPAIMSASPGEKVIMTCSASSVNYMHWFQQESGTFPKRIYDTSKLASGVPAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                   polypeptides with disulfide of prokaryotic cells in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecular chaperone; PelB signal sequence; scFvOxazolon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 444; DB 3; I
Pred. No. 4.2e-27;
7; Mismatches 15;
                                                                                                                                                                                                                  Preparation of water-soluble eukaryotic
bridges e.g. rPA, comprises cultivation
presence of arginine or amide compound.
                                                                                                                                                                                                                                                                                                                          German
                                                                                               Ţ
                                                                                               AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PelB-scFvOxazolon fusion protein.
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ilarity 80.0%;
Conservative
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99EP-00107412
                                               99EP-00107412
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N-PSDB; AAC66074.
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88; Conser
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Ħ Schwarz Schaeffner J, Rudolph R, Ambrosius D,

WPI; 2001-246712/26. N-PSDB; AAF77806.

Producing naturally folded eukaryotic proteins e.g. antibodies, interferon, hormones or proteases that contain two or several cysteines linked by disulfide bridges comprises co-expression of a molecular chaperone.

Disclosure; Page 19; 35pp; English.

The present invention relates to a method for production of a naturally folded eukaryotic protein containing two or more cysteines linked by disulfide bridges. The method comprises co-expression and secretion into the periplasm of a molecular chaperone via an expression vector coding for the chaperone. The expression vector also encodes a signal sequence. The method is useful for producing a naturally folded eukaryotic protein such as an antibody, antibody fragment, interferon, protein hormone or a protease containing two or several cysteines linked by disulfide bridges. The present sequence is a fusion protein composed of the PelB signal sequence and ScFvOxazolon. This sequence was used in the method of the present invention

Sequence 255 AA;

Length 255; Indels 27; 15; Score 444; DB 4; Pred. No. 4.2e-27; ; Mismatches 19 7; 73.3%; llarity 80.0%; Conservative Local Similarity 88; Query Match Matches Best

DIELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRIYDTSKLASGVPAR 60 |||:|||||| SKLSSGVPAR 132 DIELTQSPAIMSASPGEKVTMTCSASSSVRYMNWFQQKSGTSPKRWIYDT

191

241

0; Gaps

110 61 LSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADA

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Sequence 34, Appl
Sequence 2, Appli
Sequence 17, Appl
Sequence 251, App
Sequence 251, App
Sequence 243, App
Sequence 243, App
Sequence 241, App
Sequence 241, App
Sequence 5, Appli
Sequence 5, Appli
Sequence 5, Appli
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                                                                                                                                                                                                                                                                                         Published Applications AA:*

cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

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cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

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US-10-803-622-251

US-10-803-622-243

US-10-803-653-251

US-10-803-653-243

US-10-803-653-241

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3-09-144-886- 1S-10-803-622 1S-10-803-622 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-622 1S-10-803-622 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653 1S-10-803-653	3-037-2 52-889- 34-788- 18-856- 3-037-6 52-889-
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ALIGNMENTS

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Sequence 34, Application US/10410907A

Sequence 34, Application US/10410907A

Publication No. US20030215880A1

GENERAL INFORMATION:

APPLICANT: Bennis R. Burton

APPLICANT: R. Anthony Williamson

APPLICANT: R. Anthony Williamson

TITLE OF INVENTION: MOTIF-GRAFTED HYBRID POLYPEPTIDES AND

TITLE OF INVENTION: WOSES THEREOF

FILE REFERENCE: 22908-1229

CURRENT APPLICATION NUMBER: US/10/410,907A

CURRENT FILING DATE: 2003-04-08

FRIOR APPLICATION NUMBER: 60/371,610

PRIOR FILING DATE: 2002-04-09

NUMBER OF SEQ ID NOS: 36

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 34

LENGTH: 215

THENGTH: 215
                                                                                                                                                                                                                                                                                                                                                                                                                Score 471; DB 14;
Pred. No. 7.3e-36;
9; Mismatches 17;
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larity 77.8%;
Conservative
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ORGANISM: Artificial
FEATURE:
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Matches 91
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GENERAL INFORMATION:

APPLICANT: SWEET, RAYMOND W.

GENERAL INFORMATION:

APPLICANT: TORNETTA, WARK A.

APPLICANT: TRUNEH, ALEMSEGED

APPLICANT: WATTAM, TREVOR A.

TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

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TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

TITLE OF INVENTION: USFUL IN TREATMENT OF RANK LIGAND MEDIATED DISORDERS

TITLE OF INVENTION NUMBER: US/10/344,779

PRIOR FILING DATE: 2003-02-19

PRIOR FILING DATE: 2001-08-21

PRIOR FILING DATE: 2000-08-21

PRIOR FILING DATE: 2000-08-21
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Pred. No. 5.4e-35;
; Mismatches 16
                                                                                                                         APPLICANT: TORNETTA, MARK A.
APPLICANT: TORNETTA, MARK A.
APPLICANT: TRUNEH, ALEMSEGED
APPLICANT: TRUNEH, TREVOR A.
TITLE OF INVENTION: ANTI-RANK LIGAND MONOCLO
TITLE OF INVENTION: USEFUL IN TREATMENT OF
FILE REFERENCE: GP50033
CURRENT APPLICATION NUMBER: US/10/344,779
CURRENT FILING DATE: 2003-02-17
PRIOR FILING DATE: 2001-08-21
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-08-21
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FASTSEQ for Windows Version 4.0
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NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
                                    Sequence 2, Application US/10344779
Publication No. US20030211106A1
GENERAL INFORMATION:
APPLICANT: SWEET, RAYMOND W.
APPLICANT: TORNETTA, MARK A.
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Best Local S
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CEA, Conjugates Comprising Said Antibody, in an Adept System
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Pred. No. 1.3e-34;
3; Mismatches 19;
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                         16;
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Cambridge Antibody Technology Limited
Medical Research Council
McCafferty, John
Pope, Anthony
           ed, No. 5.4e-35
Mismatches 10
                                                                                                                                                                                                                 US-09-910-059-17;
Sequence 17, Application US/09910059;
Patent No. US20020142359A1;
GENERAL INFORMATION:
APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CE;
TITLE OF INVENTION: Their Therapeutic use in
                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 2001-07-23
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR FILING DATE: 1998-10-29
PRIOR FILING DATE: 1997-04-29
PRIOR FILING DATE: 1997-04-29
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1997-02-14
PRIOR FILING DATE: 1996-05-04
NUMBER OF SEQ ID NOS: 131
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5. US20040157214A1
              Pred
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Griffiths, Andrew
Jackson, Ronald
Holliger, Kasper
Marks, James
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ilarity 76.9%;
Conservative
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Clackson,
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Johnson,
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; Sequence 251, Applicat;
; Publication No. US2004;
; GENERAL INFORMATION:
; APPLICANT: Cambridge
; APPLICANT: Medical R
; APPLICANT: McCaffert
; APPLICANT: Pope, Ant
; APPLICANT: Hoogenboc
; APPLICANT: Hoogenboc
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90; Conser
             Similarity
89; Conser
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LENGIH: 235
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APPLICANT

29;

Length 1

DB 16;

Score 459;

75.78;

Query Match

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APPLICATION NUMBER: GB 9104744.9
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                                                                       Specific
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JANT: Bonert, Timothy
OF INVENTION: Methods for Producing Members of
REFERENCE: 13839-00013
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C: Cambridge Antibody Technology
F: Medical Research Council
F: McCafferty, John
F: Pope, Anthony
F: Johnson, Kevin
F: Hoogenboom, Hendricus
F: Griffiths, Andrew
F: Jackson, Ronald
F: Holliger, Kasper
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APPLICANT: Chiswell, David
APPLICANT: Winter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
TITLE APPLICATION NUMBER: US/10/803,622
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81.5%; Pred. No. 2e-34;
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                                                                                                                                          OR FILING DATE: 1990-07-10

OR APPLICATION NUMBER: GB 9022845.3

OR APPLICATION NUMBER: GB 9022845.3

OR APPLICATION NUMBER: GB 9022845.3

OR FILING DATE: 1990-10-19

OR APPLICATION NUMBER: GB 9024503.6

OR FILING DATE: 1990-11-12

OR APPLICATION NUMBER: GB 9104744.9

OR FILING DATE: 1991-03-06

OR APPLICATION NUMBER: GB 9110549.4
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                                                                                                                                                                                                                                                                                                    FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
FILING DATE: 1991-07-10
APPLICATION NUMBER: US 07/971,857
FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
                                                                              FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
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PatentIn version 3.1
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Chiswell, David
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CURRENT FILING DATE: 2004-0
PRIOR APPLICATION NUMBER: GE
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; Sequence 251, Application No. US201; GENERAL INFORMATION:
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NUMBER OF SEQ
SOFTWARE: Pate
SEQ ID NO 251
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Best Local S
Matches 88
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APPLICANT: Pope, Anthony
APPLICANT: Pope, Anthony
APPLICANT: Pope, Anthony
APPLICANT: Hoogenboom, Hendricus
APPLICANT: Griffiths, Andrew
APPLICANT: Jackson, Ronald
APPLICANT: Grackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Chiswell, David
APPLICANT: Chiswell, David
APPLICANT: Winter, Gregory
APPLICANT: Winter, Gregory
APPLICANT: Monethy
TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: VL of scFv from mouse immunized with 2-phenyl-5-oxazolone
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Cambridge Antibody Technology Limited
Medical Research Council
McCafferty, John
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PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-05-15
PRIOR FILING DATE: 1991-05-15
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1991-07-10
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 272
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 251
LENGTH: 108
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PRIOR APPLICATION NUMBER: GB 9015198.6
PRIOR FILING DATE: 1990-07-10
PRIOR APPLICATION NUMBER: GB 9022845.3
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APPLICATION NUMBER: GB 9022845.3
FILING DATE: 1990-10-19
APPLICATION NUMBER: GB 9024503.6
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NUMBFP
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5. US20040157214A1
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Similarity 81.5%;
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ORGANISM: Artificial
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Publication No. US200
GENERAL INFORMATION:
APPLICANT: Cambridge
APPLICANT: Medical
APPLICANT: Medical
APPLICANT: Mcaffer
APPLICANT: Mcaffer
APPLICANT: Hoogenbo
APPLICANT: Griffith
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Sequence 241, Application US/10803622
Publication No. US20040157214A1
GENERAL INFORMATION:
APPLICANT: Cambridge Antibody Techno
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ORGANISM: Artificial Sequence
FEATURE:
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Clackson, Timothy
Chiswell, David
Winter, Gregory
Bonert, Timothy
VENTION: Methods for
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272
                NUMBER OF SEQ ID NOS: 272
SOFTWARE: Patentin version 3.1
SEQ ID NO 243
LENGTH: 108
                                                                                                  Sequence
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FILING DATE: 1995-06-07 OF SEQ ID NOS: 272
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                                                                                  TYPE: PRT ORGANISM: Artificial FEATURE:
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NUMBER OF SEQ ID NOS
SOFTWARE: Patentin v
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87; Conser
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SOFTWARE: Pace
SEQ ID NO 241
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APPLICANT: McCafferty, John
APPLICANT: Pope, Anthony
APPLICANT: Johnson, Kevin
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, Andrew
APPLICANT: Griffiths, Andrew
APPLICANT: Holliger, Kasper
APPLICANT: Marks, James
APPLICANT: Clackson, Timothy
APPLICANT: Chiswell, David
APPLICANT: Winter, Gregory
APPLICANT: Womer, Gregory
APPLICANT: Womer, Timothy
TITLE REFERENCE: 13839-00013
CURRENT FILING DATE: 2004-03-18
PRICA APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRICE APPLICATION NUMBER: GB 9015198.6
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Pred. No. 4.7e-34;
3; Mismatches 13;
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Cambridge Antibody Technology Limited
Medical Research Council
McCafferty, John
Pope, Anthony
    PRIOR FILING DATE: 1991-03-06

PRIOR APPLICATION NUMBER: GB 9110549.4

PRIOR FILING DATE: 1991-05-15

PRIOR APPLICATION NUMBER: PCT/GB91/01134

PRIOR FILING DATE: 1991-07-10

PRIOR FILING DATE: 1993-01-08

PRIOR FILING DATE: 1993-01-08

PRIOR FILING DATE: 1995-06-07

NUMBER OF SEQ ID NOS: 272

SOFTWARE: Patentin version 3.1

SEQ ID NO 243
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OR APPLICATION NUMBER: GB 9022845.3

OR APPLICATION NUMBER: GB 9022845.3

OR APPLICATION NUMBER: GB 9022845.3

OR APPLICATION NUMBER: GB 9024503.6
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R: PCT/GB91/01134
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Sequence 243, Application US/10803653
Publication No. US20040157215A1
GENERAL INFORMATION:
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FILING DATE: 1993-01-08
APPLICATION NUMBER: US C
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                                                                           Length 108;
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   from mouse immunized with
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F: Cambridge Antibody Technology Limited
F: Medical Research Council
F: McCafferty, John
F: McCafferty, John
F: Pope, Anthony
F: Johnson, Kevin
F: Hoogenboom, Hendricus
F: Griffiths, Andrew
F: Jackson, Ronald
F: Holliger, Kasper
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                                                                                Score 448; DB 16;
Pred. No. 4.7e-34;
3; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: MINCEL, GLESOLY
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Me
FILE REFERENCE: 13839-00013;
CURRENT APPLICATION NUMBER: US/10/803,622
CURRENT FILING DATE: 2004-03-18
PRIOR PILING DATE: 1990-07-10
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-10-19
PRIOR FILING DATE: 1990-11-12
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-03-06
PRIOR FILING DATE: 1991-05-15
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 08/484,893
PRIOR FILING DATE: 1993-01-08
PRIOR FILING DATE: 1993-01-08
PRIOR APPLICATION NUMBER: US 08/484,893
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Philadelphia
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US-09-795-515-5
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                                                                                                                                      1 DIELTQSPAIMSASPGEKVTMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPAR
                                                                                      Gaps
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     enyl-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Marks, James
APPLICANT: Clackson, Timothy
APPLICANT: Clackson, Timothy
APPLICANT: Chiswell, David
APPLICANT: Chiswell, David
APPLICANT: Minter, Gregory
APPLICANT: Bonert, Timothy
TITLE OF INVENTION: Methods for Producing Members of Specific Bin;
FILE REFERENCE: 13839-00013
CURRENT APPLICATION NUMBER: US/10/803,653
CURRENT FILING DATE: 2004-03-18
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9022845.3
PRIOR APPLICATION NUMBER: GB 9024503.6
PRIOR APPLICATION NUMBER: GB 9104744.9
PRIOR FILING DATE: 1990-11-12
PRIOR APPLICATION NUMBER: GB 910549.4
PRIOR FILING DATE: 1991-03-06
PRIOR PILING DATE: 1991-05-15
PRIOR PRIOR APPLICATION NUMBER: GB 9110549.4
PRIOR FILING DATE: 1991-07-10
PRIOR PILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR PILING DATE: 1991-07-10
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 07/971,857
PRIOR APPLICATION NUMBER: US 07/971,857
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                                                                                     Indels
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                                                     Length
 immunized
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                                                                                    14;
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                                                 Score 446; DB 16;
Pred. No. 7.2e-34;
; Mismatches 14
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Pred. No. 7.2e-34;
7; Mismatches 14
    mouse
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                                                                                                                                                                                                                                                                                                     Sequence 241, Application US/10803653 Publication No. US20040157215A1 GENERAL INFORMATION:
scFv from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          version 3.1
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80.6%;
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ilarity 80.6%;
Conservative
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of
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OTHER INFORMATION: VL
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87; Conser
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    ; US-10-803-622-241
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ENGTH: 108
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Matches 87
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IELTQSPAIMSASPGEKVIMTCSASSSVNYMHWFQQESGTFPKRRIYDTSKLASGVPARL
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                                                       LSGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRA
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46th Floor
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Pred. No. 2.1e-33;
); Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Version
                                                                                                                                                                                                                                    APPLICANT: Adair, John R.
APPLICANT: Athwal, Diljeet S.
APPLICANT: Emtage, John S.
TITLE OF INVENTION: Humanised Antibodies
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mac)
                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,515
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0057
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/09/795,515
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FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
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US-09-795-515-5
; Sequence 5, Application US/09795515
; Publication No. US20030039645A1
; GENERAL INFORMATION:
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US-10-704-352-5
; Sequence 5, Application US/10704352
; Publication No. US20040071693A1
; GENERAL INFORMATION:
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One Liberty Place - 4
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Similarity 75.0%; Pr
87; Conservative 10;
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Athwal, Diljeet
Emtage, John S.
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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Place - 46th Floor
                                          Mackiewicz
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Pred. No. 2.1e-33;
); Mismatches 19;
                                          Kurtz Mac
46th Floor
                                                                                                                                                                                                  CURRENT APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US/10/704,352

FILING DATE: 07-Nov-2003

CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/795,515

FILING DATE: 28-FEB-2001

APPLICATION NUMBER: 08/846,658

FILING DATE: 01-MAY-1997

ATTORNEY/AGENT INFORMATION:

NAME: Trujillo, Doreen Yatko

REGISTRATION NUMBER: 35,719

REFERENCE/DOCKET NUMBER: CARP-0057

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jonn S.
Humanised Antibodies
Antibodies
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                            Floppy disk
1 PC compatible
3TEM: PC-DOS/MS-DOS
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                                         Washburn
Place - 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 5, Application US/10704071;
Publication No. US20040076627A1;
GENERAL INFORMATION:
APPLICANT: Adair, John R.
Athwal, Diljeet S.
Emtage, John S.
TITLE OF INVENTION: Humanised A.
                                                                                                                                                                                                                                                                                                                                                                                                            5) 568-3100
568-3439
Humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 235 amino acids
TYPE: amino acid
TOPOLOGY: linear
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                                           STREET: One Liberty R
CITY: Philadelphia
STATE: PA
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STREET: One Liberty
CITY: Philadelphia
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.larity 75.0%;
Conservative 10
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                                          ADDRESSEE: Woodcock
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                                                                                                COUNTRY: USA
ZIP: 19103
COMPUTER READABLE FORM:
                                                                                                                                                           IBM PC C
SYSTEM:
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MOLECULE TYPE: protein
;
SEQUENCE DESCRIPTION: 0
US-10-704-352-5
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CORRESPONDENCE ADDRESS
           NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
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                                                                                                                                               MEDIUM TYPE:
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87; Conser
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                                                                                                                                                            COMPUTER:
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OF
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US-10-704-071-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62 SGSGSGTEFTLEISRVKAEDVGVYYCQQLVEYPLTFGAGTKLELKRADAAPTVSIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             84 RGSGSGTSYSLTISGMEAEDAATYYCQQWSSNPFTFGSGTKLEINRADTAPTVSIF
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 Version
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Pred. No. 2.6e-33;
7; Mismatches 14;
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Pred. No. 2.1e-33;
0; Mismatches 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 82, Application US/09144886
; Patent No. US20020155114A1
; GENERAL INFORMATION:
; APPLICANT: Marks, James D
; APPLICANT: Amersdorfer, Peter
; TITLE OF INVENTION: Therapeutic Monoclonal An;
; TITLE OF INVENTION: Botulinum Neurotoxins
; FILE REFERENCE: 2500.117USO
; CURRENT APPLICATION NUMBER: US/09/144,886
; CURRENT FILING DATE: 1998-08-31
; NUMBER OF SEQ ID NOS: 98
                                                                                                                                     FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0057
                                                                                                                        658
                                   US/10/704,071
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                    ເນ
•••
                                                                                                                        08/846,
                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of
VL e
                                  APPLICATION NUMBER: US/1 FILING DATE: 07-Nov-2003
                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acid
TYPE: amino acid
TOPOLOGY: linear
                                                                                                       APPLICATION DATA: APPLICATION NUMBER: US
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                   CURRENT APPLICATION DATA:
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Similarity 75.0%;
87; Conservative 10
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nilarity 80.4%;
Conservative
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                                                                     CLASSIFICATION:
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ORGANISM: Artificial
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86; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NC SOFTWARE: Patentin
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LENGTH: 107
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Compugen Ltd
GenCore version (c) 1993 - 2004
          Copyright
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protein search, using sw model OM protein

; Search time 24.7042 {without alignments} 471.266 Million cell 13, 2004, 12:48:23 October Run on:

updates/sec

/TVSS

US-10-049-868A-4 635 score: Perfect

QVQLQESGPGLVAPSQSLSI.......MITAYAMDYWGQGTTV Sequence:

BLOSUM62 Gapop 10.0 , Gapext Scoring table:

0.5

283416 seqs, 96216763 residues Searched:

283416 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 2000000000 DB DB Minimum Maximum

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 79:*
1: pir1:*
2: pir2:*
1: pir3:* H 0 M 4 Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	A Property Charles 17	g neavy chain v	g neavy chain v	g neavy chain v	g neavy ch	d heavy chain V	g heavy chain V	d heavy chain v	d heavy chain or	d heavy chain v	d heavy chain v	Theater chain t	g neavy chain v	y heavy chain v	s heavy chain v	g neavy chain v	g neavy chain V	g gamma-2A cha	g heavy chain V	g heavy chain V	g heavy chai	g heavy chain V	g heavy chain V	gamma-1 chain	y heavy chain V	heavy chain pr	Theavy chain (a	Theath, chain 11	heavy chain	amma-2b chain
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79 75.4 109 2 S	79 75.4 114 2 S	.5 75.4 144 2 S	77 75.1 107 2 S	76 75.0 1 07 2 S	75 74.8 100 2 S	.5 115 2 S	71 74.2 111 2 S	71 74.2 114 2 S	70 74.0 107 2 S	.5 73.6 110 2 P	57 73.5 95 2 S	54 73.1 110 2 S	53 72.9 109 2 P	.5 72.7 115 2 82	55 71.7 101 2 S0

ALIGNMENTS

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23-Jul-199
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RESULT 1
S52446
Ig heavy chain V region precursor - mouse
C; Species: Mus musculus (house mouse)
C; Date: 08-May-1995 #sequence_revision 03-Aug-1995 #text_change 23-C; Accession: S52446
R; Berdaz, J; Kraehenbuhl, J.P.
Submitted to the EMBL Data Library, November 1994
A; Description: Specific amplification by the polymerase chain react A; Reference number: S52445
A; Accession: S52446
A; Molecule type: DNA
A; Cross-references: EMBL: X82692; NID: 9673441; PIDN: CAA58013.1; PID: C; Genetics:
A; Introns: 16/1
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 34-116/Domain: immunoglobulin homology <IMM>
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Pred. No. 3.1e-45;
2; Mismatches 10;
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Best Local Similarity 89.3%;
Matches 109; Conservative
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21-Jan-2000

2-phenyloxazolon t O response RESULT 2
S11100
Ig heavy chain V region (clone NQ2-20.5.3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-C;Accession: S11100
R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A;Title: mRNA sequences define an unusually restricted IgG response A;Reference number: S07331; MUID:83271467; PMID:6877353
A;Accession: S11100
A;Molecule type: mRNA
A;Residues: 1-112 <NAT>
C;Superfamily: immunoglobulin V region; immunoglobulin homology

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-phenyloxazolor
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                                QVQLKRSGPGLVAPSQSLSITCTVSGFSLTSSGVHWVRQPPGKGLEWLGVIWAGGSTNYN
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S55028
Ig heavy chain V region - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 23-Aug-1995 #sequence_revision 19-Oct-1995 #text_change 23-C; Date: 23-Aug-1995 #sequence_revision 19-Oct-1995 #text_change 23-C; Accession: 855028
R; Jeffrey, P.D.; Schildbach, J.F.; Chang, C.Y.; Kussie, P.H.; Margo, J. Mol. Biol. 248, 344-360, 1995
A; Title: Structure and specificity of the anti-digoxin antibody 40-A; Reference number: 855027; MUID:95257394; PMID:7739045
A; Reference number: 855027; MUID:95257394; PMID:7739045
A; Residues: 1-140 < JEF>
A; Residues: 1-140 < JEF>
A; Cross-references: EMBL:L31403; NID:g476717; PIDN:AAA38191.1; PID:C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Superfamily: immunoglobulin homology < IMM>
F; 34-116/Domain: immunoglobulin homology < IMM>
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C; Date: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_cl
C; Accession: S11101
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein,
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG
A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Accession: S11101
A; Accession: S11101
A; Status: preliminary
A; Molecule type: mRNA
C; Superfamily: immunoglobulin V region; immunoglobulin home
F; 15-97/Domain: immunoglobulin homology <IMM>
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Pred. No. 5.1e-43;
7; Mismatches 14;
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Pred. No. 6.2e-43
2; Mismatches
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l Similarity 84.3%;
102; Conservative
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Best Local Similarity 82.6%;
Matches 100; Conservative
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C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 23-Jul-1999
C; Accession: PL0087
R; Meek, K.; Hasemann, C.; Pollok, B.; Alkan, S.S.; Brait, M.; Slaoui, M.; U. Exp. Med. 169, 519-533, 1989
A; Title: Structural characterization of antiidiotypic antibodies; evidence A; Reference number: PL0080; MUID:89094248; PMID:2492056
A; Reference number: PL0080; MUID:89094248; PMID:2492056
A; Residues: 1-120 <MEB>
A; Residues: 1-120 <MEB>
A; Residues: 1-120 <MEB>
A; Residues: 1-120 <MEB>
C; Superimental source: strain BALB/C
A; Note: the sequence shown here is from the VH region of an antiidiotypic m C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-97/Domain: immunoglobulin homology <IMM>
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                                                                                                                                   OVOLKESGPGLVAPSQSLSITCTVSGXSLTSYGVHWVRQPPGKGLEWLGVIWAGGSTNYN
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                                                                                                                                                                                                                                                                                                         Ig heavy chain V region (ASWS1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1
C;Accession: S38563
R;Monestier, M.; Losman, L.J.; Novick, K.E.; Aris, J.P.
Submitted to the EMBL Data Library, September 1993
A;Description: Molecular analysis of mercury-induced anti-nucleolar anti-A;Reference number: S38563
A;Reference number: S38563
A;Accession: S38563
A;Accession: S38563
A;Kesidues: 1-117 <MON>
A;Residues: 1-117 <MON>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;15-97/Domain: immunoglobulin homology <IMM>
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Pred. No. 3.1e-43;
3; Mismatches 13;
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Pred. No. 1.6e-43;
5; Mismatches 8;
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                                               Score 536; DB 2;
Pred. No. 8.7e-44
1; Mismatches
               <IMM>
              immunoglobulin homology
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ilarity 86.7%;
Conservative
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Best Local Similarity 86.0%;
Matches 104; Conservative
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larity 85.8%;
Conservative
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103; Conser
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                ;15-97/Domain:
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Best Local S
Matches 103
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C; Species: Mus musculus (house mouse)
C; Species: Mus musculus (house mouse)
C; Date: 26-Oct-1899 #sequence_revision 31-Dec-1993 #text_change 23-Jul-1999
C; Accession: A32456
R; Dombrink-Kurtzman, M.A.; Johnson, L.S.; Riordan, G.S.; Bedzyk, W.D.; Voss Jr., E.W.
J. Biol. Chem. 264, 4513-4522, 1989
A; Title: Variable region primary structures of a high affinity anti-fluorescein immunocy
A; Reference number: A32456
A; Reference number: A32456
A; Recession: A32456
A; Status: Preliminary
A; Molecule type: mRNA
A; Residues: 1-139 < DOM>
A; Residues: 1-139 < DOM>
A; Residues: 1-139 < DOM>
A; Residues: the authors translated the codon CAC for residue 20 as Gln, and CAC for residue C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
E; 34-116/Domain: immunoglobulin homology < IMM>
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C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-Jan-C; Accession: S11103
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG response to A; Reference number: S07331; MUID:83271467; PMID:6877353
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$11108

Ig heavy chain V region (clone NQ5-78.2.6) - mouse (fragmes)
C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_c]
C; Accession: $11108
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein,
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG
A; Reference number: $07331; MUID:83271467; PMID:6877353
A; Molecule type: mRNA
A; Residues: 1-112 <NAT>
C; Superfamily: immunoglobulin V region; immunoglobulin home
F; 15-97/Domain: immunoglobulin homology <IMM>
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Pred. No. 1.3e-42;
; Mismatches 12;
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Pred. No. 2.3e-42;
2; Mismatches 9;
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Best Local Similarity 85.0%;
Matches 102; Conservative
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Best Local Similarity 84.2%;
Matches 101; Conservative
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C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-C; Accession: S11106
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG response A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Accession: S11106
A; Molecule type: mRNA
A; Residues: 1-114 <NAT>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 15-97/Domain: immunoglobulin homology <IMM>
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-C;Accession: S11102
R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
R;Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
A;Title: mRNA sequences define an unusually restricted IgG response
A;Reference number: S07331; MUID:83271467; PMID:6877353
A;Accession: S11102
A;Molecule type: mRNA
A;Residues: 1-116 <NAT>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
E;15-97/Domain: immunoglobulin homology <IMM>
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Score 527; DB 2; Le
Pred. No. 6.3e-43;
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Pred. No. 6.4e-43;
2; Mismatches 9;
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Best Local Similarity 84.3%;
Matches 102; Conservative
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larity 84.3%;
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C; Species: Mus musculus (house mouse)
C; Date: 19-Mar-1997 #sequence_revision 23-Aug-1997 #text_change 21-C; Accession: S11099
R; Kaartinen, M.; Griffiths, G.M.; Markham, A.F.; Milstein, C.
Nature 304, 320-324, 1983
A; Title: mRNA sequences define an unusually restricted IgG response A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Reference number: S07331; MUID:83271467; PMID:6877353
A; Residues: 1-114 <NAT>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 15-97/Domain: immunoglobulin homology <IMM>
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S42484
Ig heavy chain V region (4B1 VH) - mouse
C; Species: Mus musculus (house mouse)
C; Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 2.
C; Accession: S42484
R; Gilbert, D.; Brard, F.; Margaritte, C.; Delpech, A.; Tron, F. submitted to the EMBL Data Library, March 1994
A; Description: An idiotype D23-bearing polyspecific, murine anti-A; Reference number: S42484
A; Reference number: S42484
A; Reference number: S42484
A; Residues: preliminary
A; Molecule type: mRNA
A; Residues: 1-116 <GIL>
A; Cross-references: EMBL: Z30962; NID:g461325; PIDN: CAA83216.1; PIC; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
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  A;Accession: S11103
A;Molecule type: mRNA
A;Residues: 1-115 <NAT>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-97/Domain: immunoglobulin homology <IMM>
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                                                                                              Score 517; DB 2; I
Pred. No. 5.6e-42;
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8.6e-42;
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1 Similarity 82.6%; Pred. No. 8.66
100; Conservative 2; Mismatches
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ilarity 83.5%;
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[Species: Mus musculus (house mouse)

[Species: Mus musculus (house mouse)

[Species: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-C; Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-C; Accession: A49049

[R; Armandola, E.A.; Mariani, S.M.; Zwickl, M.; Hardman, N.; Ferrone, S. Eur. J. Immunol. 22, 2893-2899, 1992

[A; Title: Molecular analysis of anti-idiotypic monoclonal antibodies in A; Reference number: A49049

[A; Reference number: A49049

[A; Reference number: A49049

[A; Residues: preliminary

[A; Molecule type: nucleic acid

[A; Residues: 1-122 < ARM>

[A; Residues: 1-122 < ARM>

[A; Residues: 1-122 < ARM>

[A; Residues: nucleic acid

[A; Residues: nuclei
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C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1998 #sequence_revision 19-Mar-1998 #text_change 21-C;Accession: S26322
R;Stark, S.E.; Caton, A.J.
J. Exp. Med. 174, 613-624, 1991
A;Title: Antibodies that are specific for a single amino acid inter A;Title: Antibodies that are specific for a single amino acid inter A;Reference number: S26309; MUID:91341421; PMID:1908510
A;Reference number: S26309; MUID:91341421; PMID:1908510
A;Residues: 1-106 <STA>
A;Residues: 1-106 <STA>
A;Residues: 1-106 <STA>
A;Cross-references: EMBL:X59182
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;9-91/Domain: immunoglobulin homology <IMM>
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Pred. No. 2.2e-40;
; Mismatches 13;
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                                                          Score 507.5; DB 2
Pred. No. 4.5e-41;
; Mismatches 11
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immunoglobulin homology
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                                                          th 79.9%; Similarity 80.3%; 98; Conservative
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Gaps
Query Match
Best Local Similarity 85.1%; Pred. No. 2.1e-40;
Matches 97; Conservative 2; Mismatches 7; Indels
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Search completed: October 13, 2004, 13:00:06 Job time : 24.7042 secs

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               Copyright
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- protein search, using sw model OM protein

Run on:

updates/sec October 13, 2004, 12:47:33 ; Search time 121 Seconds (without alignments) 575.375 Million cell updat

score: Title: Perfect so Sequence:

US-10-049-868A-4 635 1 QVQLQESGPGLVAPSQSLSI......MITAYAMDYWGQGTTVTVSS 121

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

1825181 seqs, 575374646 residues Searched:

1825181 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description				מינות (נומסחפא ל	homo garie	55 camelus	aines omod	ines omod	ines omon	ines omod	homo agra-	homo sapi				33 camelus	homo sapi	nomo sapi	02 hom	homo sap	homo s	homo s	homo sapi	homo sari	ines omod	homo omor	AT POWOR SEPTE	ACTURE TO TOWN OF	acessay nome sap	d00465 camelus
SUMMARIES	ID	091X92	9NG	HV43 MOUSE	MOUS	ı	HV02_XENLA	OGGMX6	BAD00255	Q9UL73	HV3J HUMAN	Q96E <u>Y</u> 0	O6GMX1	Ö6GMX7	06P418	AAH63384		0233	ງ ລຸດ	6/52/2 08mm3))	BAC85202	CSWUX4	Q6GMX5	Q9BU10	Q96AA6	Q9HCC1	Q8N5K4	06N089	578	85	O1 XE	000465
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32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

PRT; 482 AA.	, Created) , Last sequence update) , Last annotation update)	ta; Craniata; Vertebrata; Euteleostomi; ia; Sciurognathi; Muridae; Murinae; Mus	ouse L.H., Derge J.G r L., Shenmen C.M.,	H., Schaefer ax S.I., Wan .A., Rubin G	F., Casavant ki S., Carnin	J., Malek J.A., Gun arcia A.M., Gay L.J	Noweighten E.D., Lu A., Gibbs K.A., nan M., Madan A., Rodrigues S., Sanchez J.A.C., Shevchenko Y., Bouffard G.G.,	1 J.w., Green E.D., Dickson M.C., I J., Schmutz J., Myers R.M., Butterfield . U., Smailus D.E., Schnerch A., Schein J	analysis of more than 15,000 full-length	A. 99:16899-16903(2002).		TAMBI/Combani/Jobbi datakanan								4.
PRELIMINARY;	001 (TrEMBLrel. 19, 001 (TrEMBLrel. 19, 004 (TrEMBLrel. 26,	llus (Mouse). 11us (Mouse). 1; Metazoa; Chordata; Eutheria; Rodentia; D=10090;	ROM N.A. /N; TISSUE=Cc 388257; PubMe R.L., Feingc .D., Collins	., Zeeberg , Jordan H. ., Marusina	, Soares M.) [.J., Usdin '	S., McEwan P.J., M.S., Worley K.C.,	Helton E., Kette , Madan A., Youn	r.w., louchman U.w Z.A.C., Grimwood J., ki M.I., Skalska U.,	Marra M.A.; and initial	1. Acad. Sci. U.S.A.	FROM N.A. B/N; TISSUE=Colon;	R.,	181; AAH1118	; F33932. 0; 1A7N.		PR003006; Ig_	PR003596; IG	54; Cl-set; 47; iq; l.	406; IGV;	50835; IG LIKE;
1X92 Q91X92	01-DEC-2001 01-DEC-2001 01-DEC-2001 01-MAR-2004	Mus musculus Eukaryota; M Mammalia; Eu NCBI_TaxID=1	SEQUENCE FI STRAIN=FVB, MEDLINE=22 Strausberg Klausner R	Altschul Hopkins R Diatchenk	Stapleton Brownstei Raha S.S.	Bosak S.A Richards	Fahey J., Whiting M	Rodriguez A. Krzywinski M	Generatic	Proc. Nat.	SEQUENCE FR STRAIN=FVB/	Strausberg Submitted	EMBL; BC01	HSSP, PO18	InterPro; InterPro:	InterPro;	InterPro;	Pfam; PF00	SMART; SMOO	PROSITE; E

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21-JUL-1986
21-JUL-1986
01-OCT-2004
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                                            GVINTGGSTNYN
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01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2003 (TrEMBLrel. 25, Last annotation update)
Single chain Fv (Fragment).
Mus musculus (Mouse).
Plasmid pHEN1.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Murinae; NCBI_TaxID=10090;
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SEQUENCE FROM N.A.
STRAIN=BALB/C;
MEDLINE=98169018; PubMed=9510199;
MEDLINE=98169018; PubMed=9510199;
MEDLINE=98169018; PubMed=9510199;
MEDLINE=98169018; Robert M., Baensch M., Sohns B., Arstrantsch W., Kola A., Klos A., Koehl J.;
Bautsch W., Kola A., Klos A., Koehl J.;
"Site-Directed C3a-Receptor Antibodies from Phage Display Lik"
Site-Directed C3a-Receptor Antibodies from Phage Display Lik
J. Immunol. 160:2947-2958(1998).
R MBL; RJ222590; CAA10890.1; -.
R PIR; FJ3932; FJ3932.
R InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
           82
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Pred. No. 2e-43;
3; Mismatches 16;
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10;
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Pred. No. 2.2e-44
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HV43 MOUSE
ID HV43 MOUSE STANDARD;
AC P01819;
DT 21-JUL-1986 (Rel. 01, Created)
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ilarity 79.3%;
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           Query Match
Best Local Similarity 81.3%;
Matches 100; Conservative
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
Plasmid.
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121 AA;
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Sciurognathi; Muridae; Murinae; Mus
                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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SEQUENCE FROM N.A.

MEDLINE=81012133; PubMed=6774258;

Sakano H., Maki R., Kurosawa Y., Roeder W., Tonegawa S.;

"Two types of somatic recombination are necessary for the of complete immunoglobulin heavy-chain genes.";

Nature 286:676-683(1980).

-!- MISCELLANEOUS: The sequence shown is translated from differentiated gene isolated from a myeloma that secritification of contains 1 immunoglobulin-like domain.
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(Rel. 01, Last sequence update)
(Rel. 45, Last annotation updating vegion PJ14 precursor.
21-JUL-1986 (Rel. 01, Last sequence update) 29-MAR-2004 (Rel. 43, Last annotation updat Ig heavy chain V region MOPC 141 precursor. Mus musculus (Mouse).
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Mismatches
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MEDLINE=81012133; PubMed=6774258
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Eukaryota; Metazoa; Chordata; C:
Mammalia; Eutheria; Rodentia; S:
NCBI_TaxID=10090;
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ilarity 75.8%;
Conservative
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Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE;
Immunoglobulin V region; Si
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                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-s or send an email to license@isb-sib.ch).
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kano H., Maki R., Kurosawa Y., Roeder W., Tonegawa S
wo types of somatic recombination are necessary for
complete immunoglobulin heavy-chain genes.";
ture 286:676-683(1980).
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Pred. No. 2.3e-38;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                               Signal
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                                                                                                                                                                                                                                                                                                                                                                EMBL; V00767; CAA24148.1; -. PIR; A02095; HVMS14.

PDB; 1A7N; X-ray; H=20-115.

PDB; 1A7P; X-ray; H=20-115.

PDB; 1A7P; X-ray; H=20-115.

PDB; 1A7P; X-ray; H=20-115.

PDB; 1G7H; X-ray; B=20-115.

PDB; 1G7J; X-ray; B=20-
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larity 88.5%;
Conservative
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85; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                   HV45 MOUSE
P01821;
21-JUL-1986
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SEQUENCE
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"Two ty
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Matches 8
                                        Nature
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HV45_MOUSE
  ID
AC
DT
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MEDLINE=82075900; PubMed=6273429;
Kataoka T., Nikaido T., Miyata T., Moriwaki K., Honjo T.,
"The nucleotide sequences of rearranged and germline immunoglobulin typenes of a mouse myeloma MC101 and evolution of VH genes in mouse.";
genes of a mouse myeloma MC101 and evolution of VH genes in mouse.";
J. Biol. Chem. 257:277-285(1982).
-!- SIMILARITY: Contains 1 immunoglobulin-like domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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MEDLINE=88176921; PubMed=2451244;
Schwager J., Mikoryak C.A., Steiner L.A.;
Schwager J., Mikoryak C.A., Steiner L.A.;
"Amino acid sequence of heavy chain from Xenopus laevis IgM deduced from cDNA sequence: implications for evolution of immunoglobulin domains.";
Proc. Natl. Acad. Sci. U.S.A. 85:2245-2249(1988).
21-JUL-1986 (Rel. 01, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Ig heavy chain V region MC101 precursor.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P20957;
01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Ig heavy chain V region XIG14 precursor (Fragment).
Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
NCB1_TaxID=8355;
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Ig-like.
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF00047; ig; 1. SMART; SM00406; IGv; 1. PROSITE; PS50835; IG LIKE; 1. Immunoglobulin V region; Signal
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HSSP; P01820; 1G7J.
InterPro; IPR007110; Ig-like
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 83.5%;
81; Conservative
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PIR; A02096; GIMS10.
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116
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116
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Query Match
Best Local S
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                      Jh a collaboration EMBL outstation
                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Primary B-Cells;

X MEDLINE=22388257; PubMed=12477932;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buerow K.H., Schaefer C.F., Bhat N.K.,

Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Both Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Robards S.A., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Kahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Whiting M., Madan A., Young A.C., Schwerz R.M., Butterfield Y.S.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B.,

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    64 MSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDYWGQGTTVTVSS 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -length human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 LOESGPGLVAPSQSLSITCTVSGISLNRYGVHWVROPPGKGLEWLGVIWTGGSTNYNSAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                            Ig heavy chain V region XIG14
Ig-like.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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    SIMILARITY: Contains 1 immunoglobulin-like domain.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 401.5; DB 1;
Pred. No. 4.9e-34;
19; Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      465 AA.
                                                                                                                                                           EMBL; J03632; AAA49791.1; ~.
PIR; B31933; B31933.
HSSP; P01820; 1A7N.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin V region; Signal.
NON TER 1
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128
135
15080 MW;
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ilarity 61.9%;
Conservative 1
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Best Local Similarity
Matches 73; Conser
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SEQUENCE
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Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus
NCBI_TaxID=9838;
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AB091893; BAD00255.1; ".
                                                                                                             EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
TISSUE=Peripheral blood;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies reflecting camel
                                                                                                                                                                                                                                                                                                                                                                                                              B3A9B7D0FDB1386E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunoglobulin heavy chain VHDJ region (Fragment).
              99:16899-16903 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.3%; Score 389.5; DB 2;
larity 62.8%; Pred. No. 3.8e-32;
Conservative 17; Mismatches 23;
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58.1%; Pred. No. 5.2e-32;
iive 20; Mismatches 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (Arabian camel)
                                                                                Strausberg R.;
Strausberg R.;
Submitted (JUN-2004) to the EMBL/Ger.
EMBL; BC073766; AAH73766.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR003110; Ig-like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003006; Ig_W.
Pfam; PF07654; Cl-set; 3.
Pfam; PF07047; ig; 4.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Camelus dromedarius (Dromedary)
                   U.S.A.
sequences."
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                                                        SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                    Sci.
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75; Conserv
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129 AA;
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           Proc. Natl. Acad. [2]
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ID BAD00255
AC BAD00255;
DT 02-MAR-2004 (
DT 02-MAR-2004 (
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Q96EY0;
01-DEC-2001
01-DEC-2001
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RGVYSMDYWG
  IITAYAMDYWG
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                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                            stomi;
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         S.M.,
                                                                                                                                                            Euteleo
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                                                                                                                                    region
                                                                                                                                                                                                                                                                                                                                                      119
                                                                                                                                                                   Homo.
                                                                                                                                                                                                       Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M. Young D.C.; "Myosin-reactive autoantibodies in rheumatic carditis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Ig heavy chain V-III region HIL.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Butelé
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                       Indel
                                                                                                                                                                                                                                                                                                                                                     2; Length
                                                                                                                                                                                                                                                                                                                                     1BDB86B6420EA0BE CRC64;
                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                       O9UL73;
O9UL73;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2004 (TrEMBLrel. 26, Last annotation update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
YNSALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDR
                                                                                                                                                                                                                                                                                                                                                                      29;
                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Willing B., Van der Merwe P.L., Kalis N.N.,
                                                                                                                                                                                                                                           87:184-192(1998)
                                                                                                                                                                                                                                                                                                                                                  59.2%; Score 376; DB 2. larity 62.0%; Pred. No. 2e-31; Conservative 15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                      Clin. Immunol. Immunopathol. 87
EMBL; AF035041; AAD56277.1; -.
PIR; PH0876; PH0876.
PIR; S12416; S12416.
HSSP; P01820; 1G7J.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                    13219 MW;
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                                                                                                                                                                                                                                                                                               SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
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                                121
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                                                                                                                                                  Homo sapiens (Human)
                                         : | | | | : | | KGTLVTISS
                                OGTTVTVSS
                                                                                                                                                                                                                                                                                                                                                           l Similarity
75; Conserv
                                                                                                                                                                                                                                                                                                                                   119 AA;
                                                                                                                                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Euthe:
NCBI_TaxID=9606
                                                                                                                                                                                                                                                                                                                                                                                                                                                    121
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                                                                                                                                        (Fragment)
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59
                61
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NON TER
SEQUENCE
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                                                                                                                                                                                                                                  fetus."
                                                                                                                                                                                                                                                                                                                                                            Local
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Matches
                                                                        RESULT
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119
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TISSUE=Primary B-Cells;

X MEDLINE=22388257; Pubmed=12477932;

A trausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 NSALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDYWGQGTTVTV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OVOLQESGPGLVAPSQSLSITCTVSGISLNRYGVHWVRQPPGKGLEWLGVIWTGGS-TNY
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Catarrhini; Hominidae; Homo.
                                                                                                                                                myeloma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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MEDLINE=79124695; PubMed=420800;
Chiu Y.-Y.H., Lopez de Castro J.A., Poljak R.J.;
"Amino acid sequence of the VH region of human myeloma
cryoimmunoglobulin IgG Hil.";
Biochemistry 18:553-560(1979).
-!- MISCELLANEOUS: This chain was isolated from an IgG:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CRC64;
                                                                                                                                                                                                Contains 1 immunoglobulin-like domain
                                                                                                                                                                                                                                                                                                                                                                                  InterPro; Lencis 1.

Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG LIKE; 1.

Direct protein sequencing; Immunoglobulin V region;

Direct protein sequencing; Immunoglobulin V region;

Pyrrolidone carboxylic acid.

1 112 Ig-like.

Pyrrolidone carboxylic
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Last sequence update)
Last annotation update)
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27;
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larity 54.9%; Pred. No. 4.2e-31;
Conservative 26; Mismatches 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           620 AA
                                                                                                                                                                                       -!- SIMILARITY: Contains 1 immunoglobulir PIR; A02054; G1HUHL.
HSSP; P01772; 2FB4.
G0; G0:0005576; C:extracellular; NAS.
G0; G0:0006953; F:antigen binding; NAS.
G0; G0:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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13566 MW;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
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(TrEMBLrel. 19, 1
(TrEMBLrel. 26, 1
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121 AA;
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67; Conserv
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Dickson M.C

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and mouse
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Best Local
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                                                                                  Proc.
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Q6GMX7
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                            length human
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X TISSUE-Spleen;

X TISSUE-Spleen;

X TISSUE-Spleen;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Grouse L.H., Schaefer G.D.,

X Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Rosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Rahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                           1 OVOLOESGPGLVAPSOSLSITCTVSGISLNRYGVHWVROPPGKGLEWLGVIWTGGSTNYN
                                                                                                                                                                                                                                                                                                                                                                                                                                         27 QVQLQESGPGLVKPSETLSLTCTVSGGSISSYYWSWIRQPAGKGLEWIGRIYTSGSTNYN
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Hulyk S.W.,
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   Skalska U., Smailus D.E., Schnerch A., Schein
                                                                                                                                                                                                                                                                                                                                                                       620;
Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hômo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                           databases
                                                                                                                                                                                                                                                                                                                                             CRC64;
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27, Last sequence update)
27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                       Score 371.5; DB 2;
Pred. No. 4.1e-30;
19; Mismatches 28;
                                                                                                                                                                                                                                                                                                                                            990A1A4A6E8FF27B
                                                                                                                           the EMBL/GenBank/DDBJ
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                                                                                                           Strausberg R.;
Submitted (JUL-2001) to the EMBL
EMBL; BC011857; AAH11857.2; -.
PIR; S15590; S15590.
HSSP; P01820; 1G7J.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig.cl.
InterPro; IPR003596; Ig.w.
Pfam; PF00047; ig; 4.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGW; 1.
PROSITE; PS50835; IG LIKE; 5.
PROSITE; PS00290; IG MHC; UNKNOW SEQUENCE 620 AA; 68125 MW; 5
                                                                                                                                                                                                                                                                                                                                                                                                   19;
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60.3%;
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Q6GMX1;
Q6GMX1;
05-JUL-2004 (TrEMBLrel. 2'05-JUL-2004 (TremBLrel).
                                                                                                                                                                                                                                                                                                                                                                                      Similarity 60... 3; Conservative
                                                                                   SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   146
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Matches
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P SEQUENCE FROM N.A.

TISSUE=Primary B-Cells;

X TISSUE=Primary B-Cells;

X Farausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Itschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22
28
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                                                                                             analysis of more than 15,000 full-length human
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                                Butterfield
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                                                         Schein
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                                                         A.,
                                                                                                                                                                                                                                                   databases
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YNSALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARD---
Green E.D., Dickers R.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                           U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          57.6%; Score 366; DB 2;
larity 60.5%; Pred. No. 1.1e-29;
Conservative 14; Mismatches 27
                                                                                                                                                                                                                                                to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   477 AA
                                           Д., Ж.
          Blakesley R.W., Touchman J.W., Green Rodriguez A.C., Grimwood J., Schmutz Krzywinski M.I., Skalska U., Smailus Jones S.J., Marra M.A.; "Generation and initial analysis of m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
                                                                                                                                                                                                                        Strausberg R.;
Strausberg R.;
Submitted (JUN-2004) to the EMBL/Gen
EMBL; BC073773; AAH73773.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig.al.
InterPro; IPR003597; Ig.al.
InterPro; IPR003596; Ig.al.
InterPro; IPR003596; Ig.al.
InterPro; IPR003596; Ig.al.
Efam; PF07654; Cl-set; 3.
Pfam; PF00047; ig; 4.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
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27,
27,
                                                                                                                            cDNA sequences.
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Homo sapiens (Human).
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Mammalia; Eutheria;
NCBI_TaxID=9606;
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                                                                                                                                                Natl. Acad.
                                                                                                                                                                                        FROM N.A
                                                                                                                                                                                        SEQUENCE FROM
TISSUE=Spleen;
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Hypothetical
SEQUENCE 47
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05-JUL-2004
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X MEDLINE=22388257; PubMed=12477932;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Heieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heieh F.,

RA Brownstein M.J., Usdin T.B., Former A.A., Rubin G.M., Hong L.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Bosak S.A., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
                                                                                                      ength human
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atne P.H.,
Hulyk S.W.,
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Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y. Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E. Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length hu
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CRC64;
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Last annotation update)
                                                                                                                sequences.";
ad. Sci. U.S.A. 99:16899-16903(2002)
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Pred. No. 1.6e-29;
7; Mismatches 29;
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InterPro; IPR003599; IG.
InterPro; IPR003597; IG-like.
InterPro; IPR003597; IG-like.
InterPro; IPR003597; IG-like.
InterPro; IPR003597; IG-like.
InterPro; IPR003596; IG-like.
InterPro; IPR003596; IG-like.
Pfam; PF07654; Cl-set; Z.
Pfam; PF00047; ig; 3.
SMART; SM00409; IG; 4.
SMART; SM00406; IG; 4.
PROSITE; PS50835; IG-LIKE; 4.
PROSITE; PS00290; IG-MHC; UNKNOWN Hypothetical protein.
SEQUENCE 477 AA: F.
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larity 59.5%;
Conservative
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(TrEMBLrel.
(TrEMBLrel.
                                                                                                                                                                                               B-Cells;
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                                                                                                                                                                                                             Strausberg R.;
Submitted (JUN-2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 72; Conser
                                                                                                                                    Proc. Natl. Acad.
                                                                                                                                                                            FROM N
                                                                                                                    CDNA
                                                                                                                                                                                              TISSUE=Primary
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Name=IGHD;
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5-JUL-2004
5-JUL-2004
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X MEDLINE=22388257; PubMed=12477932;

X MEDLINE=22388257; PubMed=12477932;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

X Strausberg R.L., Feingold E.A., Rapher C.M., Schuler G.D.,

A lischul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
                                                                                                                                                                                                                                                                                                                                                  of more than 15,000 full-length human
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Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E. Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length be a second for the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and initial analysis of more than 15,000 full-length because the contraction and the contracti
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Strausberg R.;
Submitted (DEC-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BC063384; AAH63384.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig.cl.
InterPro; IPR003596; Ig.MHC.
InterPro; IPR003596; Ig.V.
Pfam; PF07654; Cl-set; 3.
Pfam; PF07654; Cl-set; 3.
SMART; SM00409; IG; 1.
SMART; SM004009; IG: 1.
SMART; SM00406; IGV; 1.
PROSITE; PS00290; IG LIKE; 4.
PROSITE; PS00290; IG MHC; UNKNOWN 2.
SEQUENCE 576 AA; 63363 MW; FBB97C949D720F1E CRC64;
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27, Last sequence update)
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larity 62.3%; Pred. No. 2.1e-29;
Conservative 14; Mismatches 29;
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AAH63384;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence up
02-MAR-2004 (TrEMBLrel. 27, Last annotation
Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vert
Mammalia; Eutheria; Primates; Catarrhini; Ho
                                                                                                                                                                                                                                                                                                                                                                                         and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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Fahey J., Helton D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Whiting M., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
Jones S.J., Marra M.A.;
T. "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
T. "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
T. "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
T. "Generation and initial analysis of more than 15,000 full-length human broc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).

T. Schein J. "
Schein 
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Best Local Similarity 62.3%; Pred. No. 2.1e-29;
Matches 76; Conservative 14; Mismatches 29;
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GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                Copyright
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protein search, using sw model 1 OM protein

Seconds 2004, 12:44:23 October 13, Run on:

updates/sec ; Search time 112.933 (without alignments) 384.353 Million cell

US-10-049-868A-4 635 score:

QVQLQESGPGLVAPSQSLSI......MITAYAMDYWGQGTTVTVSS Title: Perfect sc Sequence:

121

BLOSUM62 Gapop 10.0 , Scoring table:

0.5 Gapext

2002273 segs, 358729299 residues Searched:

2002273 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 2000000000 Minimum DB Maximum DB

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Geneseq_23Sep04:*
geneseqp1980s:*
geneseqp2000s:*
geneseqp2001s:*
geneseqp2002s:*
geneseqp2003as:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ALIGNMENTS

RESI	LT 1		
AAR	AAR21268 TD bab01068 standard: wrotoin, 112 am		
X	initiated scandard, process; its AA.		
AC	AAR21268;		
×			
DJ	21-MAY-1992 (first entry)		
×			
DΕ	Murine VH group 2 chain E specific for phox.		
×			
Κ	Fd; bacteriophage; gene III; filamentous: phagemid: gansid: goat: milus:	. m.	2
Ϋ́	93p; binding; adsorption; gene VIII: diverse repertoire.	111111111111	3
Ϋ́	specific binding pairs: replicable genetic display package		
×	Converse to the second of the		
SO	Synthetic.		
×			
II L	Voy		

Ξ Location/Qualifiers
31. .35
/label= CDR1
50. .65
/label= CDR2
98. .102
/label= CDR3
/note= " D/N-X-G-X-X motif" Key Binding-site Binding-site Binding-site

90GB-00015198. 90GB-00022845. 90GB-00024503. 91GB-00004744. 90GB-00015198 10-JUL-1990; 19-OCT-1990; 12-NOV-1990; 06-MAR~1991; 15-MAY-1991; 10-JUL-1990; WO9201047-A 23-JAN-1992

(CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY (MEDI-) MED RES COUNCIL.

Griffiths AD; Hoogenboom HRJ, KS, Johnson , Marks J, Pope AR, Holliger KP, Mccafferty J Jackson RH,

WPI; 1992-056862/07

Producing members of specific binding pairs - by expression in recombinant host cells with a secreting replicable genetic display package.

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a single chain Fv library from an immunised mouse. The library produces a clive VH sequence is one of antibody fragments specific for 2-phenyl-5-caiverse repertoire of antibody from minned from mRNA from mice immunised with phox coupled to chicked serum albumin. The VH and VL kappa sequences were sequenced sequences were sequenced revealing the phage surface as fusions with gene III. The resulting library of clones was diverse. Twenty three hapten binding clones were sequenced revealing the eight different VH genes (A-H) in a variety of pairings with seven different VK genes (a-g) (see AAR21264-92). Nearly all the VH genes belonged to gp 1, with only cone, "E", being of gp 2 (VHOx1). Of the twenty three clones sequenced, only one was of type "E". Most of the clones were Vk-d combinations. The kd of VH-B/Vk-d for phox-GABA was 10 nM. Only two other combinations (of eleven tested) were found to have higher values. This suggests that phage bearing scFv fragments having weak affinities can be selected with antigen, probably due to the avidity of the multiple antibody heads on the phage. See also AAR21260-307, 309-311; AAR22450, 565-581
                                                                                                                                                                                                                                                        is suggests that phage selected with antibody heads on 565-581
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                                           (AAR21264-71) found to be
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                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                           Score 557; DB 2; I
Pred. No. 4.8e-44;
); Mismatches 5;
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               24; 209pp;
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                                                                                                                                                                                                                                                                                                                                                                                           09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ablating/killing aberrant prostate specific membrane antigen-expressing
                                                                                     mouse anti
                                                                               This sequence represents a heavy chain variable region from a mouse ant: Vascular Adhesion Protein-1 (VAP-1) antibody. This sequence is included in the specification for comparison with the antibody of the invention. The murine antibody sequences of the invention may be used in the production of a chimeric mouse-human anti-VAP-1 antibody. The nucleic acid molecules, polypeptides or antibodies are useful in treating VAP-1 mediated inflammatory disorders, such as rheumatoid arthritis, inflammatory bowel disease, autoimmune diseases or psoriasis. The chimeric VAP-1 antibody is further used for in vitro and in vivo diagnostic applications, including in vivo immunoscintigraphic imaging inflammation sites. The chimeric MAD's of the invention have improved kinetic properties compared to the corresponding murine antibodies.
and
                                                                                                                                                                                                                                                                                                                                                             SALMSRLSISKDNSKSOVFLKMNSLOTDDTAMYYCARDRSTMITAYAMDYWGOGTTVTVS
                                                                                                                                                                                                                                                                                                                                                                                                                                        New chimeric anti-Vascular Adhesion Protein-1 monoclonal antibodies encoding nucleic acid molecules, useful for diagnosing and treating chronic inflammatory disorders, e.g. rheumatoid arthritis or psorias
                                                                                                                                                                                                                                                                                                                                      9
                                                                                                                                                                                                                                                                                                        Length 115;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Murine variable heavy chain amino acid sequence MuVHIB
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 555; DB 8; I
Pred. No. 7.5e-44;
1; Mismatches 5;
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                                                            English.
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2002US-0362612P.
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                                                                                                                                                                                                                                                                                                         87.4%;
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                                                                                                                                                                                                                                                                                Sequence 115 AA;
                                                            Fig
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08-MAR-2002;
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                                                              Disclosure;
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The present invention describes a method (MI) for ablating or killing an aberrant prostate specific membrane antigen (PSMA) expressing cell (e.g. an epidermal and adermal cell). MI comprises contacting the cell, or a vascular endothelial cell). MI comprises contacting the cell, or a vascular endothelial cell proximate to the cell, with an antibody (or its antigen-binding fragment), which binds specifically to the extracellular domain of PSMA in an amount sufficient to ablate or kill the cell. The antihoflammatory and antialleristic, antiarthritic, dermatological, cytostatic, antiinflammatory and antialleristic activities, and can be used in vaccines. MI is useful for treating a skin disorder in a subject, by administering to the subject, an amount of an antibody which binds specifically to the extracellular domain of PSMA (the subject is a dministering to the subject, an amount of an antibody which binds specifically to the extracellular domain of PSMA (the subject is a dministeribly human and is having, or at risk of, a skin disorder.). The skin disorder is a dermal or an epidermal disorder, and is selected from psoriasis (preferably chronic stationary psoriasis, psoriasis contasis, and localised pustular psoriasis, psoriatic arthritis, exfoliative dermatities, pityriasis rubra plans, lichen nitidus, christionare dermatities, pityriasis rubra plans, lichen plans, lichen nitidus, inflammatory or neoplastic disorder of the epidermal or neglecancerous lesion. MI is also useful to treat or prevent disorders involving aberrant activity of PSMA and ABR44613 to ABR44613 to ABR44613 represent sequences used in the exemplification of the beaucerous and in the exemplification.
                                specific
         vith
      contacting the cell was ar domain of prostate
cells for treating skin disorders, by contacting antibody that binds to the extracellular domain
                                                                                                         Disclosure, Fig 12B; 225pp; English.
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                                                        antigen.
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116 120 58 9 QVQLQESGPGLVAPSQSLSITCTVSGISLNRYGVH--WVRQPPGKGLEWLGVIWTGGSTN AMDYWGQGTS Gaps 4. Length 125; Indels .. 60 , DB 6; 1.3e-43; Score 553; DB Pred. No. 1.3e-3; Mismatches 3; 87.1%; ilarity 87.2%; Conservative Similarity 125 121 Sequence 125 AA VTVSS ||||| VTVSS 109; Query Match Best Local S Matches 109 Н S O 61 117 g ò В à ö

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ScFv; anti-human seminal plasma protein;
                                                                                                                          monoclonal antibody E4B7; mouse; murine; human; carboxypepti
fusion protein; antibody-directed enzyme prodrug therapy; AI
methotrexate-alpha-peptide prodrug; prostate cancer; cytosta
                                                                                        Anti-human seminal plasma protein ScFv/hCPA fusion protein.
                        667
                      standard; protein;
                                                                     entry)
                                                                                                             Single chain antibody;
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RESULT 4
ABP97414
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                      in single chain antibody E4B7
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                   "Anti-human seminal plasma protein single ch
ly derived from murine monoclonal antibody E4
regions; targets fusion protein to prostate
                                                                                 138. .245
/note= "Corresponds to monoclonal antibody E4B7
chain variable region"
                                                                                                                                                               activates
                                                                                                                                                             carboxypeptidase A;
prodrug"
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               SCFV
                                        1. .122
/note= "Corresponds to mcchain variable region"
123. .137
/label= ScFv lint-
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/label= ScFv/hCPA_linker
247
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Location/Qualifiers
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                                  and VL regions; cells"
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/label= hCPA
/note= "Human o
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Protein
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Pharmaceutical kits for use in the treatment of prostate cancer, com anti-human seminal plasma protein single-chain antibody/human carboxypeptidase fusion protein and a prodrug of methotrexate-alphapeptides

29pp; Chinese. 4 Fig Claim 3;

The invention relates to a pharmaceutical kit for antibody-directed enzyme prodrug therapy (ADEPT) to treat prostate cancer. The kit comprises several respective containers for holding anti-human seminal plasma protein single-chain antibody (ScFv)/human carboxypeptidase A (hCPA) fusion protein (ABP97414), a prodrug of methotrexate-alpha-peptide and a pharmaceutically-acceptable auxiliary for medication. The invention also encompasses the anti-human seminal plasma protein ScFv/hCPA fusion protein, and the nucleic acid encoding it (ABZ75116). This was constructed by linking DNA encoding the ScFv (itself generated by linking DNA encoding the heavy and light chain variable regions (VH and VL) of the murine anti-human seminal plasma protein monoclonal antibody E4B7) to DNAs encoding human carboxypeptidase A via a linker. The ScFv portion of the fusion protein enables it to be specifically targeted to prostate cancer cells, where the hCPA portion can then activate the prodrug which in turn has cytostatic activity towards the cancer cells. The invention permits the fusion protein to be easily produced in large quantities for use in ADEPT for prostate cancer treatment. The present sequence

Seguence

idase A; hCPA; ADEPT; atic; gene;

Homo sapiens

Synthetic. Chimeric.

Length 667; DB 6; 3e-42; Score 547; Pred. No. 86.1%; 86.8%; Query Match Best Local Similarity insulin-dependent vulgaris and spresent the NKG2D

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disease; tumour;
lon; liver; ovary;
uterus; cervix;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; NKG2D; NKG2D receptor complex; cancer; infectious disease; tumour; autoimmune disease; head; neck; stomach; oesophagus; colon; liver; ovary; intrahepatic bile duct; pancreas; lung; larynx; breast; uterus; cervix; prostate; kidney; testis; thyroid; bladder; brain; melanoma; myeloma; Fv; sarcoma; leukaemia; lymphoma; virus; bacterium; fungus; protozoan; DAP10; helminth; cytostatic; antimicrobial; immunomodulatory; 11B2D10; 6H7E7; 8G7C10; 6E5A7; 11B2D10x4-7; 8G7C10x4-7; 6E5A7x4-7; P4-2; P4-3; P4-14; P5-15; P5-2; P5-3; P5-9; P5-10; P5-11; P5-23; 3B10xP4-3; 3B10xP4-14; P53 tetramerisation domain; 3B10xP5-2; 3B10xP5-23.
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treating
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Hofmeister R;
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N-PSDB; AAS97141.
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oesophagus, colon, liver, intrahepatic bile ducts, pancreas, lung, larynx, breast, ovary, uterus, cervix, prostate, kidney, testis, thyroid, bladder or brain, or a melanoma, myeloma, sarcoma, leukaemia or lymphoma. The infectious diseases can be caused by viruses, bacteria, fungi, protozoa or helminths. The autoimmune diseases include multiple sclerosis, Grave's disease, ankylosing spondylitis, acute anterior

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The present sequence represents the heavy chain variable region of murine monoclonal antibody 2C4. This antibody binds to human sialoadhesin factor -2 (SAF-2). The antibody is useful for treating or preventing allergic rhinitis, allergies, asthma, anemia, eczema or diseases such as lymphoma, leukemia or systemic mastocytosis in a mammal. It is also useful for detecting the presence of a cell, especially eosinophil in a sample, by detecting binding of the antibody to SAF-2. The antibody can be coupled to toxins, antiproliferative drugs or radionuclides to kill cells in areas of excessive SAF-2 expression
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                                                                                                                                                                                           Monoclonal antibody 2C4; sialoadhesin factor-2; SAF-2; allergic rlallergy; asthma; anemia; eczema; lymphoma; systemic mastocytosis; leukemia; eosinophil.
                                                                                                                    4.
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uveitis, Goodpasture's syndrome, myasthenia gravis, insulindiabetes mellitus, rheumatoid arthritis, pemphigus vulgaris autoimmune hepatitis. Sequences AAU72820-AAU72875 represent receptor and the polypeptides of the invention
                                                                                           Length 253;
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                                                                                 , DB 5; Lt
1.2e-42;
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ilarity 88.5%;
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N-PSDB; AAQ37459.
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11-JUN-1992;
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19-JUN-1993
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                                                            cell populations, partic. hybridomas - to link or more non-contiguous DNA sequences to facility
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/label= Vkappa_NQ2/12.4
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/label= VH_NQ2/12.4
114. .115
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/note= "Linker |
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a forward Ck primer nested in respect to the primer used for cDNA. The assembled product within the cells is then amplified with nested primers annealing to the 5' end of the VH gene and the 3' end of the Jk segment. In-cell PCR may be used to determine gene linkage analysis, particularly for the cloning of gene combinations that are polymorphic within a population of cells, such as the rearranged genes for Ig or TCR V regions. (Updated on 25-MAR-2003 to correct PN field.)
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128. .235
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larity 86.0%;
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    The sequences given in AAR32840-43 show the mature heavy chain VH domains and the Vk light chain genes of the antiphenyloxazolone hybridomas NQ2/12.4 and NQ10/12.5 which have been linked via a linker peptide by incell PCR. The cDNA encoding these peptides was synthesised using forward primers annealing to the Ck gene and the JH segment, followed by assembly with linker primer of the Ck grimers based on the VH3 leader sequence and a forward Ck primer nested in respect to the primer used for cDNA. The assembled product within the cells is then amplified with nested primers annealing to the 5' end of the VH gene and the 3' end of the Jk segment. In-cell PCR may be used to determine gene linkage analysis, particularly for the cloning of gene combinations that are polymorphic within a population of cells, such as the rearranged genes for Ig or TCR V regions. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           anti-CTLA4;
antibody;
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                                                                                                                                                                                                                                                                                      GVIWTGGSTNYN
                                                                                                                                                                                                                                                                                                                                             SALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDYWGQGTTVTVS
                                                                                                                                                                                                                                                                                                                                                                       -YWGQGTLVTVS
                                                                                                                                                                                                                                                                                                                  GVIWAGGSTNYN
                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody-toxic group conjugate comprising an antibody that
lizes a molecule expressed only on activated T cells, usefu
ting immune response for treating autoimmune disorder, all
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                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                          Score 538; DB 2; I
Pred. No. 6.3e-42;
; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26 heavy chain variable region
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Tsurushita
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larity 86.0%;
Conservative
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N-PSDB; AAH76438.
                                                                                                                                                                                                                                               l Similarity
104; Conser
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                                                                                                                                                                                                          235 AA;
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modulating
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The invention relates to an antibody-toxic group conjugate comprising an antibody that specifically recognises a molecule expressed only on activated T cells, and a toxic group. The T cell molecule is preferably human cytotoxic T lymphocyte associated antigen 4 (CTLA4). The antibody of the invention is a humanised anti-CTLA4 antibody comprising a sequence of 128 or 142 amino acids fully defined in the specification. The antibody-toxic group conjugate is useful for modulating the immune consistency immune ersponse to a graft, allergic response or an autoimmune disorder, immune response to a graft, allergic response or an immune response to a therapeutic protein. The antibody is also useful for T-cell typing, for isolating specific IL-creeptor-bearing cells or fragments of the receptor, for vaccine preparation, and for determining the effectiveness of an agent to downergulate CTLA4 activity. The present sequence is the heavy chain variable region of mouse antibody 26. It was used in the construction of the humanised anti-CTLA4 antibody of the invention
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                                                                                                                                                                                                                                                                                                                                                           ; melanoma;
cell carcinoma;
                                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                            Length 142;
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/note= "residues Xaa are unspecified"
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                                                                                                                                                                                                                                                                        2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chain variable region consensus sequence.
                                                                                                                                                                                                                                                             , DB 4;
4.5e-42;
                                                                                                                                                                                                                                                           Score 537; DB Pred. No. 4.5e-6; Mismatches
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ilarity 84.6%;
Conservative
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neuroblastoma; glioma; soft
tumor-associated antigen.
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                                                                                                                                                                                                                                      Sequence 142 AA;
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and polypeptide(s)

and related polynucleotide(s)

Monoclonal antibody 1A7

7; Page 84; 123pp; English

response

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sequences. The MAD 1A7 and polypeptides can be used for eliciting an anti-GD2 immune response. The polypeptides can also be used for detecting or purifying anti-GD2 antibody. The products can be used for treating GD2 -associated diseases, e.g. melanoma, neuroblastoma, glioma, soft tissue carcinoma, and small cell carcinoma. They can be used for palliating the disease or for reducing the risk of recurrence. The present sequence represents the heavy chain variable region consensus sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Delaying recurrence and/or development of ganglioside GD2-associated
tumor in individual, by administering antibody containing light and heavy
chain variable region sequences contained in sequence of specified amino
                                                                               response in
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iciting an ar
r detecting (
      melanoma
                                                                                                                                                                                                                                                                                    CTVSGISLNRYGVHWVRQPPGKGLEWLGVIWTGGSTNYN
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                                                     The invention provides a monoclonal antibouy (rem., elicits an anti-GD2 (tumor-associated antigen) immunologica humans. MAb 1A7 has defined light and heavy chain variable sequences. The MAb 1A7 and polypeptides can be used for eli-GD2 immune response. The polypeptides can also be used for the fining anti-GD2 antibody. The products can be used for the inventage of the form of the second contages.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               heavy chain variable region; mouse; ganglioside GD2-associa;
melanoma; antibody; anti-idiotype; monoclonal antibody; 1A7
neuroblastoma; glioma; sarcoma; small cell lung cancer.
                                                                                                                                                                                                                                    Length 119
                                                                                                                                                                                                                                                            Indels
  GD2-associated disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          resembling 1A7 #7.
                                                                                                                                                                                                                                 Score 535; DB 2; I
Pred. No. 5.6e-42;
; Mismatches 12;
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96US-00591196.
96US-00752844.
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Local Similarity 86.0%;
Les 104; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody heavy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
                                     Disclosure, Fig 3C; 74pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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16-JAN-1996;
21-NOV-1996;
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   to
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 useful
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Matches
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Fig 3; 82pp; English.

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Example

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        The invention relates to the recurrence and/or development of a ganglioside GD2-associated tumour, e.g. melanoma, in an individual which is delayed by administration of an antibody comprising light and heavy chain variable region sequences of the anti-idiotype monoclonal antibody 1A7. The antibody is used for delaying recurrence and/or development of GD2-associated tumour, e.g. melanoma, neuroblastoma, glioma, sarcoma, or small cell lung cancer, in individual, and for treating individual with GD2-associated tumour. The present sequence is a unique peptide region from an antibody sequence resembling the anti-idiotype antibody 1A7 heavy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse; monoclonal antibody; 1A7; anti-idiotype antibody; ganglioside GD; Cytostatic; melanoma; neuroblastoma; small cell lung cancer; tumour; antibody; variable region; VH; VL; glioma; soft tissue sarcoma; vaccine
                                                                                                                                                                                                                    QVQLQESGPGLVAPSQSLSITCTVSGISLNRYGVHWVRQPPGKGLEWLGVIWTGGSTNYN
                                                                                                                                                                                                                                                                 OVQLKESGPGLVAPSQSLSITCTVSGFSLTSYGVHWVRQPPGKGLEWLGVIWGDGSTNYN
                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel anti-idiotypic monoclonal antibody 1A7, that is capable of recruiting a tumor-specific response against glycosphingolipid GD2, useful for treating a GD2-associated disease e.g., melanoma, glioma,
                                                                                                                                                                                                ..
7
                                                                                                                                                                       Length 119;
                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibody VH region consensus sequence.
                                                                                                                                                                    Score 535; DB 6; I
Pred. No. 5.6e-42;
; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SK;
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96US-00591196.
99US-00293533.
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                                                                                                                                                                    Similarity 86.0%;
4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   standard; protein;
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16-JAN-1996;
15-APR-1999;
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The invention relates to a monoclonal antibody 1A7 (an anti-idiotype antibody eliciting an anti-GD2 (ganglioside) response). Also included are an antibody producing cell deposited under ATCC Accession No. HB-11786 (or its progeny), a polynucleotide comprising a sequence encoding a comprises at least 5 consecutive amino acids from a variable region of comprises at least 5 consecutive amino acids from a variable region of consecutive nucleotides that is capable of forming a stable duplex with a polynucleotide encoding the light or heavy chain variable region of polynucleotide consisting of a variable region encoding sequence polynucleotide consisting of a variable region encoding sequence polynucleotide, a fusion polypeptide comprising 1A7, a humanised antibody comprising 5 consecutive amino acids from the 1A7 variable regions and a vaccine comprising the antibodies. The antibodies are useful for eliciting an immune response in an individual, and for treating a GD2-associated diseases is chosen from melanoma, neuroblastoma, goft tissue sarcoma, and small cell carcinoma. The individual has a clinically detectable tumour, and the method is for treating the GD2-associated diseases. 1A7 is and the method is for treating a tumour that was previously detected in the individual and individual at the contract of individual at the contract of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    se. 1A7 is
ly detected in
etectable at the
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sectable tumour,
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antibody are
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                      $8555555555555555555555555555555x&
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120 118 9 9 SALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDYWGQGTTVTVS QVQLQESGPGLVAPSQSLSITCTVSGISLNRYGVHWVRQPPGKGLEWLGVIWTGGSTNYN GVINGDGSTNYN 2 9. Length 11 Indels ; DB 7; 5.6e-42; Mismatches Score 535; Pred. No. 3 84.3%; illarity 86.0%; Conservative Similarity Sequence 119 AA; 119 S 121 · W 104; 61 61 Н 121 119 Query Match Best Local Best Loc Matches g ВВ ò à d

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ody; 11f8 group;
                                                                                        Heavy chain; variable region; anti-DNA; monoclonal; antibody; 1. hairpin; diagnosis; inflammatory glomerulonephritis; systemic lupus erythematosus; screening; treatment; prevention;
                                                                       Anti-DNA antibody 11f8 group heavy chain variable region.
                                                                                                                                                                                                                 II
                                                                                                                                                         Location/Qualifiers
                AAW07438 standard; protein; 123 AA
                                                                                                                                                                                                                  framework
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36. .49
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                                                                                                                                       Synthetic.
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The present sequence is the heavy chain variable region of the group 11f8 putative consensus anti-DNA monoclonal antibody (MAb), which has a high affinity for single stranded DNA, low or no affinity for double stranded DNA and specifically binds a DNA hairpin. The MAb can be used diagnose disorders associated with the pathological complexation of DNA, e.g. inflammatory glomerulonephritis and systemic lupus erythematosus. It can also be used to generate reagents to screen for pharmaceutical agents, and treat and/or prevent an above disorder. The sequence was derived by aligning homologous anti-DNA MAb, whose sequences have been published, as well as several MAb of other specificities obtained from a database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NSALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRS-TMITAYAMDYWGQGTTVT
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Pred. No. 7.2e-42
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Mismatches
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scAb;

single chain; antibody; recombinant;

Fusion protein encoded by Ox VH-hinge-VL insert

(revised) (first en

25-MAR-2003 08-JUN-1993

AAR32569;

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e; secretable; Bos taurus.

r peptide; linker; Bc

Spacer rDNA;]

Synthetic.

269 AA.

AAR32569 standard; protein;

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ies (scabs) and other
functional proteins
data derived from
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                                                                                                                                                                                                                                            hinge-VL insert which was used as part of a method for cloning secretable, biologically active single chain antibodies (scAbs) secretable fusion proteins having at least 2 distinct functional or domains. NOTE: This patent has been indexed using data derive patent WO9302198-A. (First Major Country Equivalent). (Updated 2003 to correct PF field.)
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                                                         /note= "CBHI]
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                                                                                                                                                                                                                                                                                                                                        80
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                                                                                                                        anti-death receptor 4 antibodies useful for treating cancer: related disorders such as rheumatoid arthritis, sjogren's me, Grave's disease and diabetes mellitus.
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Pred. No. 5e-41;
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larity 81.0%;
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updates/sec ; Search time 87.2208
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447.653 Million cell

US-10-049-868A-4 632 score: Title: Perfect

.. MITAYAMDYWGQGTT QVQLQESGPGLVAPSQSLSI.. Sequence:

TVSS 121

Gapext BLOSUM62 Gapop 10.0 Scoring table:

residues 1356558 seqs, 322682953 Searched:

0.5

1356558 satisfying chosen parameters: of hits number Total

length: 0 length: 2000000000 seg seg DB DB Minimum Maximum

Post-processing:

Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Published Database

Published Applications AA:*

cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

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SUMMARIES

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ID	US-10-803-622-21	US-10-803-653-2	US-10-239-656-63	US-09-772-103-4	US-10-153-401	US-10-660-128-1	US-08-973-518-4	-007-093-	US-10-428-754	US-10-232-187-	US-10-239-656-	US-10-144-644	US-10-700-740-1	US-09-824-286-3
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quence 252, Ap	equence 254. Ap	equence 268, Ap	equence 108, Ap	equence 187. Ap	equence 187. Ap	equence 183, Ap	equence 183, Ap	equence 112, Ap	equence 100, Ap	quence 31, Ap	equence 31, App	equence 61, App	quence 37, Appl	Sequence 13, Ar	quence 3, Appli	Seguence 3, Appl	equence 52, App	ence 4, App	equence 16, App	equence 12, App	equence 2, Appl	quence 4, Appli	equence 5, Appl	equence 68, App	equence 53, App	equence 59, App	equence 61, App	equence 63, App	equence 65, App	equence 67,
4 US-10-207-655-25	4 US-10-207-655-25	4 US-10-207-655-2	4 US-10-169-351-10	6 US-10-803-622-18	6 US-10-803-653-18	6 US-10-803-622-18	6 US-10-803-653-18	4 US-10-194-975-11	4 US-10-194-975-10		7 US-10-351-748-3	4 US-10-059-261-6	US-08-779-784-37	5 US-10-353-721-	US-09-730-374	6 US-10-704-206-	1 US-09-842-776	5 US-10-435-299-4	4 US-10-056-052-1	4 US-10-056-052-	3 US-10-140-555-2	US-09-881-823	6 US-10-697-399-	4 US-10-010-729-6	5 US-10-682-845-5	5 US-10-682-845-5	5 US-10-682-845-6	5 US-10-682-8	5 US-10-682-845-6	5 US-10-682-845-6
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ALIGNMENTS

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APPLICANT: uniswell, bavid

APPLICANT: Winter, Gregory

APPLICANT: Winter, Gregory

ITILE OF INVENTION: Methods for Producing Members of Specific Binding Pairs

FILE REFERENCE: 13839-00013

CURRENT APPLICATION NUMBER: US/10/803,622

CURRENT FILING DATE: 2004-03-18

PRIOR FILING DATE: 1990-07-10

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-10-19

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12

PRIOR FILING DATE: 1990-11-12
                                                           T: Cambridge Antibody Technology
T: Cambridge Antibody Technology Limited
T: Medical Research Council
T: McCafferty, John
T: McCafferty, John
T: McCafferty, John
T: McCafferty, Anthony
T: Johnson, Kevin
T: Hoogenboom, Hendricus
T: Griffiths, Andrew
T: Griffiths, Andrew
T: Griffiths, Andrew
T: Griffiths, Limothy
T: Clackson, Timothy
T: Chiswell, David
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FILING DATE: 1991-05-15
APPLICATION NUMBER: PCT/GB91/01134
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1990-10-19

NUMBER: GB 9024503.6

1990-11-12

NUMBER: GB 9104744.9

1991-03-06

NUMBER: GB 9110549.4
Sequence 218, Application US/10803622
Publication No. US20040157214A1
GENERAL INFORMATION:
                                                          APPLICANT:
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US-10-239-656-63
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Best Local S
Matches 108
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CANT: Bonert, Timothy
OF INVENTION: Methods for Producing Members of Specif
REFERENCE: 13839-00013
                                                                                                                                                                                                                                     QVQLQESGPGLVAPSQSLSITCTVSGFSLTSYGVHWVRQPPGKGLEWL
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Cambridge Antibody Technology Limited
Medical Research Council
McCafferty, John
Pope, Anthony
Hoogenboom, Hendricus
Griffiths, Andrew
Jackson, Ronald
                                                                                                                                    OTHER INFORMATION: VH of scFv from mouse immunized 0-803-622-218
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2e-46;
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Pred. No. 2e-4
0; Mismatches
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R: PCT/GB91/01134
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       APPLICATION NUMBER: US 07/971,857
FILING DATE: 1993-01-08
APPLICATION NUMBER: US 08/484,893
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NUMBER: GB 9
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APPLICATION NUMBER: PCT/
FILING DATE: 1991-07-10
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Clackson, Timothy
Chiswell, David
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                                      FILING DATE: 1995-06-07
R OF SEQ ID NOS: 272
ARE: Patentin version 3.
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ilarity 89.3%;
Conservative
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1991-07-10
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CURRENT FILING DATE: 2004-(
PRIOR APPLICATION NUMBER: GR
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APPLICATION NUMBER:
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FILING DATE: 1991-03
APPLICATION NUMBER:
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                                                                                                     TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                   Query Match
Best Local Similarity
Matches 108; Conser
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FILING DATE:
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ENGTH: 113
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APPLICANT:
TITLE OF IN
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                                                                                                                    scFv from mouse immunized with 2-phenyl-5
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                                                                                                                                                                                                                                                                                                         Score 546; DB 15;
Pred. No. 5.7e-45;
2; Mismatches 8;
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                                                                                                                                                                  DB 16;
                                                                                                                                                                Score 557; DB 1
Pred. No. 2e-46;
); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KUFER, PETER
APPLICANT: RIETHMULLER, GERT
APPLICANT: LUTTERBUSE, RALF
APPLICANT: LUTTERBUSE, RALF
APPLICANT: BORSCHERT, KATRIN
APPLICANT: BORSCHERT, ROMAN
APPLICANT: HOFMEISTER, ROBERT
TITLE OF INVENTION: MULTIFUNCTIONAL POLYPEPT
TITLE OF INVENTION: TO AN EPITOPE OF THE NR
FILE REFERENCE: 029976/0106
CURRENT APPLICATION NUMBER: US/10/239,656
CURRENT FILING DATE: 2003-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2000-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 63, Application US/10239656
Publication No. US20040038339A1
GENERAL INFORMATION:
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ORGANIȘM: Artificial Seguence
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88.5%;
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1995-06-07
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Best Local Similarity 88.9
Matches 108; Conservative
                                version
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SOFTWARE: Patentin Ver.
SEQ ID NO 63
PRIOR FILING DATE: 1995
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin vers
SEQ ID NO 218
LENGTH: 113
                                                                                                     ; FEATURE:
; OTHER INFORMATION: VH
US-10-803-653-218
                                                                                         ORGANISM: Artificial
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108; Conser
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Publication No.
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LENGTH: 476
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APPLICANT: Chatterjee, Malaya
Foon, Kenneth A.
Chatterjee, Sunil K.
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TREATMENT OF MELANOMA AND SMALL CELL CARCINOMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 SALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCAR--DRSTMITAYAMDYWGQGTTVT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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APPLICANT: Turner, Katherine
APPLICANT: Turner, Katherine
APPLICANT: Collins, Mary
APPLICANT: Gray, Gary S.
APPLICANT: Morris, Domna
APPLICANT: Hinton, Paul
APPLICANT: Hinton, Paul
APPLICANT: Tsurushita, Naoya
TITLE OF INVENTION: ANTIBODIES AGAINST CTLA4 AND USES THEREFOR
TITLE OF INVENTION: ANTIBODIES AGAINST CTLA4 AND USES THEREFOR
CURRENT APPLICATION NUMBER: US/09/772,103
CURRENT FILING DATE: 2000-01-01-23
PRIOR APPLICATION NUMBER: 60/178,473
PRIOR FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.0
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ZIP: 94304-1018

ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
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Pred. No. 2.3e-44;
; Mismatches 11
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CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16, Application US/10153401
Publication No. US20030114398A1
GENERAL INFORMATION:
APPLICANT: Chatterjee, Malaya
                                                                                                Sequence 4, Application US/09772103
Publication No. US20020039581A1
GENERAL INFORMATION:
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ilarity 84.6%;
Conservative
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Best Local Similarity
Matches 104; Conser
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US-10-153-401-16
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Pred. No. 2.9e-44;
3; Mismatches 12;
                                                                                                                                                                                                              NAME: Catherine M. Polizzi
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 304142000202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
                                                                         APPLICATION NUMBER: US 09/293,533
FILING DATE: 1999-04-15
APPLICATION NUMBER: US 08/372,676
                                                                                                                                   FILING DATE: 1995-01-17
APPLICATION NUMBER: US 08/591,196
FILING DATE: 1996-01-16
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US/10/153,401
FILING DATE: 27-Aug-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi
APPLICANT: Chuntharapai, Anan
APPLICANT: Chuntharapai, Anan
APPLICANT: Dodge, Kelly
TITLE OF INVENTION: DR4 Antibodies and Uses;
FILE REFERENCE: P1245R1P2B
CURRENT APPLICATION NUMBER: US/10/660,128
CURRENT FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: US 09/322,875
PRIOR FILING DATE: 1999-05-28
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1998-01-26
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Uses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
OTHER INFORMATION: Sequence is synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
JULE TYPE: peptide
INCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Subl. 5
-10-660-128-12
Sequence 12, Application US/10660128
Perioration No. US20040120947A1
                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 119 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 494-
TELEX: 706141
INFORMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 86.0%;
Matches 104; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                        PRIOR APPLICATION DATA
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QVQLKESGPGLVAPSQSLSITCTVSGFSLTSYGVHWVRQPPGKGLEWLGVIWAGGSINYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24,973
R: 1038-765
                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
Patentin Release #1.0,
                                                                                                                                                                        Sequence 4, Application US/09007093
Patent No. US20020025315A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IONE: (416) 595-1155

NX: (416) 595-1163

NN FOR SEQ ID NO: 4:

CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
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103; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: Patent
                                                                                                                                                                                                                                                                                                                                                            CITY: Toronto
STATE: Ontario
COUNTRY: Canada
ZIP: MSG 1R7
                                                                                                                                                                                                                                                                                                                                               Suite
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Best Local S
Matches 103
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                                                                                                               Gaps
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                                                                                                                                         VOLOESGPGLVAPSOSLSITCTVSGISLNRYGVHWVRQPPGKGLEWLG
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                                                                                       Length
                                             glutamic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Tolontario
STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,518
TILING DATE: 07-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
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Pred. No. 7.6e-44;
4; Mismatches 13;
                                                                                     Score 532; DB 16;
Pred. No. 2.7e-43;
1; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                ANTIBODIES FOR DEL
TO SELECTED CELLS
                                                                                                                                                                                             ALMSRISISKDNSKSQVFLKMNSLQTDDTAMYYCARD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sim & McBurney
1 Floor, 330 University Avenue
                    ; NAME/KEY: Misc_feature
; LOCATION: 20
; OTHER INFORMATION: Xaa may be glutamine or
US-10-660-128-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1038-781
                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/08973518
Publication No. US20010014327A1
GENERAL INFORMATION:
APPLICANT: Anand, Naveen N
                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Barber, Brian H
APPLICANT: Cates, George A
APPLICANT: Caterini, Judith E
APPLICANT: Klein, Michel H
TITLE OF INVENTION: CHIMERIC AI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.
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                                                                                                                11;
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TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 85.1%;
Matches 103; Conservative
                                                                                      83.8%;
ilarity 81.0%;
Conservative 1
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NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      140 amino acids
amino acid
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CORRESPONDENCE ADDRESS
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                                                                                                     1 Similarity
102; Conser
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STREET: 6th Fl
CITY: Toronto
STATE: Ontario
                                                                                                                                                                                                                                                                 SVTVSS
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US-08-973-518-4
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                                                                                        Query Match
Best Local S
Matches 102
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            FEATURE:
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US-08-9
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                                                                                                                                                                                                                                                                  OF
                                                                                                                                                                                                                                                                              THE
                                                                                                                                                                                                                                                                  FOR DELIVERY
                                                                                                                                                                                                                                                                              OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 531.5; DB 9;
Pred. No. 7.6e-44;
; Mismatches 13;
                                                                                                                                                                                                                                                                                CELLS
                                                                                                                                                                                            APPLICANT: Anand, Naveen N
APPLICANT: Barber, Brian H
APPLICANT: Cates, George A
APPLICANT: Caterini, Judith E
APPLICANT: Klein, Michel H
TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR
TITLE OF INVENTION: ANTIGENS TO SELECTED CE
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                     701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/007,093
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: US 08/483,576
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            140
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                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/428,754
FILING DATE: 05-May-2003
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 10/428,754
FILING DATE: 05-May-2003
                                                                                                                               CHIMERIC ANTIBODIES FOR DELIVERY (ANTIGENS TO SELECTED CELLS OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
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tte 701, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 14;
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Pred. No. 7.6e-44;
4; Mismatches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-1065
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5) 595-1155
595-1163
                                                                                                                                                                                                                                                                           Floppy disk
                      Sequence 4, Application US/10428754
Publication No. US20030228304A1
GENERAL INFORMATION:
APPLICANT: Anand, Naveen N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 140 amino acids TYPE: amino acid
                                                                                                       Judith
                                                               Naveen N
, Brian H
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                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                       Cates, George A
Caterini, Judit
Klein, Michel F
                                                                   Barber, bil.
                                                                                                                                                                                                                  STATE: Ontario
COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS
                                                                                                                                                  NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & MC
STREET: Suite 701,
                                                                                                                                                            20
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TOPOLOGY: linear
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US-10-428-754-4
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103; Conser
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           US-10-428-754-4
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                                                                                                                               TITLE
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 SALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDYWGQGTTVTVS | SALMSRLSISKDNSKSQVFLKINSLQTDDTALYYCARDGSSPY-YYSMEYWGQGTSVTVS | SALMSRLSISKDNSKSQVFLKINSLQTDDTALYYCARDGSSPY-YYSMEYWGQGTSVTVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DE THE NKG2D RECEPTOR COMPLEX
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(7)
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                                                                                      Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 531; DB 14;
Pred. No. 7.1e-44;
7; Mismatches 8;
APPLICANT: Erickson-Miller, Connie L.
APPLICANT: Kikly, Kristine K.
APPLICANT: Schleimer, Robert
APPLICANT: Schleimer, Robert
APPLICANT: Nulku, Turkan E.
TITLE OF INVENTION: Sialoadhesin Factor-2 Ar
FILE REFERENCE: GH50042-1
CURRENT APPLICATION NUMBER: US/10/232,187
CURRENT FILING DATE: 2002-08-29
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2001-03-05
PRIOR FILING DATE: 2002-01-18
PRIOR FILING DATE: 2002-01-18
PRIOR FILING DATE: 2002-01-18
PRIOR FILING DATE: 2002-01-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KUFER, PETER

APPLICANT: RIETHMULLER, GERT
APPLICANT: LUTTERBUSE, RALF
APPLICANT: LUTTERBUSE, RALF
APPLICANT: LUTTERBUSE, RALF
APPLICANT: MAYER, MONIKA
APPLICANT: MAYER, MONIKA
APPLICANT: HOFMEISTER, ROBERT
TITLE OF INVENTION: TO AN EPITOPE OF THE NK
FILE REFERENCE: 029976/0106
CURRENT APPLICATION NUMBER: US/10/239,656
CURRENT FILING DATE: 2003-03-06
PRIOR APPLICATION NUMBER: EP 00106467.4
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 92
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2002-07-10
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/10239656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch 83.6%;
l Similarity 86.0%;
104; Conservative
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US-10-232-187-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23, Applica Publication No. US20 GENERAL INFORMATION:
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TYPE: PRT
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Matches 104
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LENGTH: 11
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SEQ ID NO
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic Anti-NKG2D

Sequence 2, Application US/10232187 Publication No. US20030092091A1 GENERAL INFORMATION: APPLICANT: Abrahamson, Julie A.

US-10-232-187-2

Abrahamson, Julie

rapb.

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TOPOLOGY: linear MOLECULE TYPE: protein
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      ; TOPOLOGY:
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US-10-144-644-15
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                                                  Score 526; DB 15;
Pred. No. 2.2e-43;
1; Mismatches 14;
      OTHER INFORMATION: hybridoma 6H7E7 variable heavy
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APPLICATION NUMBER: US 08/470110
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/467420
FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION: INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 19406-0939

ZIP: 19406-0939

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/144,644
FILING DATE: 2002-08-08
CLASSIFICATION:
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REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                        US-10-144-644-15

US-10-144-644-15

Sequence 15, Application US/10144644

Publication No. US20030059429A1

GENERAL INFORMATION:

APPLICANT: Ames, Robert S.

APPLICANT: Appelbaum, Edward R.

APPLICANT: Cook, Richard M.

APPLICANT: Gross, Mitchell S.

APPLICANT: Holmes, Stephen D.

APPLICANT: Theisen, Timothy W.

TITLE OF INVENTION: Recombinant IL5

TITLE OF INVENTION: Treatment of IL5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Cc
STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
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FORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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illarity 86.0%;
Conservative
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STRANDEDNESS: single
                                                                        Similarity
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INFORMATION FOR
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US-10-239-656-23
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Appelbaum, Edward R.
Chaiken, Irwin M.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
McMillan, Lynette J.
Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful i
Length 119
                                       Indels
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Score 523; DB 14;
Pred. No. 4.2e-43;
; Mismatches 12
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/700,740
FILING DATE: 04-Nov-2003
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2002-08-08

FILING DATE: 2002-08-08

APPLICATION NUMBER: US 08/470110

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/467420

FILING DATE: 06-JUN-1995

APPLICATION NUMBER: US 08/363131

FILING DATE: 23-DEC-1994

ATTORNEY/AGENT INFORMATION:

NAME: Sutton, Jeffrey A.

REGISTRATION NUMBER: 34,028

REFERENCE/DOCKET NUMBER: P50282-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

ADDRESSEE: SmithKline Beecham Corp
STREET: P.O. Box 1539-UW2220

CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             US-10-700-740-15; Application US/10700740; Sequence 15, Application US/10700740; Publication No. US20040156850A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
N FOR SEQ ID NO: 15:
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Match
Local Similarity 82.6%;
Les 100; Conservative
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Benjamin, Christopher D
Hession, Catherine A
Whitty, Adrian
TITLE OF INVENTION: COMMON GAMMA CHAIN BLOCKING AGENTS
CORRESPONDENCE ADDRESS:
ADDRESSEE: Biogen, Inc.
STREET: 14 Cambridge Center
CITY: Cambridge
STATE: Massachusetts
                                                                                      Length 119
                                                                                                                                                                                                SALMSRLSISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMDY
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12;
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ed. No. 4.2e-43;
Mismatches 12;
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/824,286
FILING DATE: 02-Apr-2001
CLASSIFICATION SATA:
APPLICATION NUMBER: 60/017,466
FILING DATE: 10-MAY-1996
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan, Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A006 PCT CIF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617 679-2838
INFORMATION FOR SEQ ID NO: 3:
COUNTY OF SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; FRAGMENT TYPE: internal
; SEQUENCE DESCRIPTION: SEQ ID NO: 3
US-09-824-286-3
                                        SEQ ID NO:
                                                                                     Score
Pred.
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; Sequence 3, Application US/09824286
; Patent No. US20020028202A1
; GENERAL INFORMATION:
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LENGTH: 112 amino acids
TYPE: amino acid
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single
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ilarity 82.6%;
Conservative
          TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: (
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TYPE: internal
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STRANDEDNESS:
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Matches 100; Conser
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                                      LQESGPGLVAPSQSLSITCTVSGFSLTSYGVHWVRQPPGKGLEWLGVIWAGGSTNYNSAL
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US-10-207-655-252
; Sequence 252, Application US/10207655
; Publication No. US20030118592A1
; GENERAL INFORMATION:
; APPLICANT: Ledbetter, Jeffrey A.
; APPLICANT: Hayden-Ledbetter, Martha S.
; TITLE OF INVENTION: BINDING DOMAIN-IMMUNOGLOBULIN FUSION PROTEINS
; FILE REFERENCE: 390069-401C1
; CURRENT FILING DATE: 2002-07-25
; NUMBER OF SEQ ID NOS: 426
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 252
: LENGTH: 121
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larity 82.6%; Pred. No. 6.7e-43;
Conservative 7; Mismatches 14;
 Mismatches
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Conservative
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uence 38, Appluence 16, Appluence 16, Appluence 16, Appliuence 4, Appliuence 4, Appliuence 67, Appluence 3, Appliuence 3, Appliuence 3, Appliuence 3, Appliuence 14, Appliuence 13, Appliuence 13, Appliuence 15, Appliu ence 67 Sequence 6. Sequence 3. Sequence 3. Sequence 1. Sequence 1. Sequence 4. Sequence 4. Sequence 2. Sequence 3. Sequence 2. Sequence 2. Sequence 1. Sequence 2. Sequence 1. Sequence 2. Sequence 3. Sequence 2. Sequence 3. Sequen iption equence equence equence equence ednence Descr: 00000000000 US-08-190-199A-67 US-08-190-199A-61 US-08-652-558-38 US-08-752-844-16 US-08-752-844-16 US-08-293-533-16 US-08-943-136-4 US-08-943-136-4 US-08-943-136-4 US-08-973-518-4 5455030-15 US-08-667-769A-15 US-08-667-769A-15 US-09-189-129-3 US-09-189-129-3 US-09-824-286-3 US-09-824-286-3 US-09-189-129-3 US-08-860-174A-2 US-08-860-174A-2 US-08-860-174A-2 US-08-860-174A-2 US-08-860-174A-2 US-08-860-174A-1 US-08-860-174A-1 US-08-861-751A-4 US-08-861-751A-4 US-08-37-411-4 US-08-37-411-4 US-08-467-420A-15 US-08-470-110A-15 US-08-940-371-15 ũ SUMMARI Π Ω Length Query Match \$\text{SQC} Score Result No.

Sequence 66, Appl	, Appl	equence 4. Appl	equence 4, Appl	equence 4, Appl	equence 4. Appl	equence 4, A	equence 66. App	equence 66. App	equence 8, Appl	equence 5, Appl	equence 12. App	equence 12, App	12, App	equence 12, App	equence 4, Appl	equence 14. App	equence 6
8-881-	06-293-	S	1 - 19		-09-2	-09-324-191-	-08-752-844-6	-09-293-533-	-08-652-558-8	-09-254-189-	2-546-1	-08-764-938-	2 - 1	US-09-131-053A-12	-08-564-1	US-07-942-245-14	0-7
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ALIGNMENTS

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US-08-19-199A-67

Sequence 67, Application US/08190199A

Patent No. 5830663

GENERAL INFORMATION:

APPLICANT: EMBLETON, Michael J.

APPLICANT: GORGCHOV, GUY

APPLICANT: JONES, Peter T.

APPLICANT: WINTER, Gregory P.

TITLE OF INVENTION: TREATMENT OF CELL POPULATIONS

NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:

ADDRESSEE: PILLSBURY MADISON & SUTRO, L.L.P.

STREET: 1100 New York Avenue, N.W.

CITY: Washington

STATE: D.C.

COMPUTER: D.C.

COMPUTER: BM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM:

MEDIUM TYPE: Flopy disk

COMPUTER: Microsoft Word

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/190,199A

FILING DATE: 13-JUL-1994

CLASSIFICATION NUMBER: PCT/GB92/01483

FILING DATE: DATE: DATE: PATAIL APPLICATION DATE: POPALICATION DATE: PCT/GB92/01483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 10-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9212419.7
FILING DATE: 11-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER: GB 9117352
10-AUG-1991
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FILING DATE: 10-AUG-1
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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US-08-190-199A-67
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RESULT 1
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80 Length 222; ; DB 2; 6.9e-48; ; Score 538; DB; Pred. No. 6.9e 2; Mismatches Query Match Best Local Similarity 86.0%; Matches 104; Conservative

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QVQLQESGPGLVAPSQSLSITCTVSGISLNRYGVHWVRQPPGKGLEWLGVIWTGGSTNYN

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                      YWGQGTTVTVS
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                    SALMSRISISKDNSKSQVFLKMNSLQTDDTAMYYCARDRSTMITAYAMD
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                                                                                                                                                                                                                           CELL POPULATIONS
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Pred. No. 7.4e-48;
; Mismatches
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                                                                                                                                                                                                                                                           SUTRO,
                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/190,199A
FILING DATE: 13-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/GB92/01483
FILING DATE: 10-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9212419.7
                                                                                                                                                                                                                                                                                Z
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                                                                                                                                                                                        APPLICANT: GOROCHOV, GUY
APPLICANT: JONES, Peter T.
APPLICANT: WINTER, Gregory P.
TITLE OF INVENTION: TREATMENT OF CE
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSEE: PILLSBURY MADISON & SI
STREET: 1100 New York Avenue, N.V.
                                                                                                                                             61, Application US/08190199A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 11-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9212419.
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                                                                                                                                                                                Michael
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Best Local Similarity 86.0%;
Matches 104; Conservative
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FORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
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amino acid
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                                                                                                                                         Sequence 61, Appliance;
Batent No. 5830663;
GENERAL INFORMATION:
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TOPOLOGY:
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US-08-190-199A-61
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CITY: Wa
STATE: D
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Sequence 16, Application US/08752844

Sequence 16, Application US/08752844

Patent No. 5935821

GENERAL INFORMATION:

APPLICANT: Chatterjee, Malaya

APPLICANT: Chatterjee, MonocLonal Antibody 1A7 AND USE FOR THE

TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CELL CARCINOMA

TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CELL CARCINOMA

NUMBER OF SEQUENCES: 66

CORRESPONDENCE ADDRESS:

ADDRESSEE: MORRISON & FOERSTER

STREET: 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 120;
            TITLE OF INVENTION: HUMANIZED ANTIBODIES AND USES TITLE OF INVENTION: THEREOF NUMBER OF SEQUENCES: 52 CORRESPONDENCE ADDRESS: ADDRESSEE: BANNED . ...
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nes 10;
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Pred. No. 5.
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,558
FILING DATE: JUNE 6, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/1B94/00387
FILING DATE: NOVEMBER 21, 1994
ATTORNEY/AGENT INFORMATION:
NAME: YANKWICH, LEON R.
REGISTRATION NUMBER: 30,237
REFERENCE/DOCKET NUMBER: 95,497-L
TELEPHONE: 617-345-9100
TELEPHONE: 617-345-9101
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                PCT/IB94/00387
                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette,
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
Sequence 38, Application US/08652558
Patent No. 5861155
GENERAL INFORMATION:
APPLICANT: LIN, AUGUSTINE YEE-TH
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illarity 86.0%;
Conservative
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104; Conser
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CITY: BOSTON
STATE: MASSACE
COUNTRY: USA
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TOPOLOGY: 111
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Best Local S
Matches 104
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; TOPOLOGY: lir
; MOLECULE TYPE:
US-08-591-196-16
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Best Local S
Matches 104
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; Patent No. 5977316
; GENERAL INFORMATION:
; APPLICANT: Chatterjee, Malaya
APPLICANT: Chatterjee, Sunil K.
; TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
; TITLE OF INVENTION: TREATMENT OF MELANOMA AND SMALL CAI
; TITLE OF SEQUENCES: 57
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 PAGE MILL ROAD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 119;
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                                                                                                                       Version #1.30
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 535; DB 2;
Pred. No. 6.6e-48;
3; Mismatches 1:
                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/752,844
FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30414-20002.21
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/08/591,196
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86.0%;
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                                                                                                                                                                                                                                                                                  TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            peptide
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FILING DATE: 16-JA
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TOPOLOGY: linear
MOLECULE TYPE: pept
                          RY: USA
94304-1018
                                                                                                                                                                                                                                                                                                                                                                amino acid
PALO ALTO
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104; Conser
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US-08-591-196-16
                            COUNTRY:
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Best Local
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STATE:
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RESULT 6
US-09-293-533-16

US-09-293-533-16

Sequence 16, Application US/09293533

Patent No. 6509016

APPLICANT: Chatterjee, Malaya
APPLICANT: Chatterjee, Malaya
APPLICANT: Chatterjee, Sunil K.

TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION: MONOCLONAL ANTIBODY 1A7 AND USE FOR THE
TITLE OF INVENTION:
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ZIP: 94304-1018

ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/293,533

FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION:

APPLICATION NUMBER: US/08/752,844
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Pred. No. 6.6e-48
3; Mismatches 11
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FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 30414-20002.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
"""" (415) 494-0792
                                                                                                                                    40,253
3R: 30414-20002.20
A24

NAME: Schiff, J. Michael
REGISTRATION NUMBER: 40,253
REFERENCE/DOCKET NUMBER: 3041
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acid
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                                                                                                                                                               Patent No. 6291208
GENERAL INFORMATION:
APPLICANT: Anand, Naveen N
APPLICANT: Barber, Brian H
APPLICANT: Cates, George A
APPLICANT: Caterini, Judith E
APPLICANT: Klein, Michel H
TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR DELIVERY OF
TITLE OF INVENTION: ANTIGENS TO SELECTED CELLS OF THE IMMUNE SYSTEM
                                                                                                                                        Gaps
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Сі
                                                                                                                Length 119
                                                                                                                                        Indels
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                                                                                                             Score 535; DB 4;
Pred. No. 6.6e-48
3; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/943,136
FILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/483,576
FILING DATE: 07-JUNE-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-733
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MBER: US 08/483,576
07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24,973
R: 1038-733
                                                                                                                                                                                                                                                                                                                                                             Sequence 4, Application US/08943136
Patent No. 6291208
                                                                                                                                        3;
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TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                            84.3%;
iilarity 86.0%;
Conservative
16
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          CHARACTERISTICS:
119 amino acids
amino acid
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SEQ ID NO:
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COUNTRY: Canada
ZIP: MSG 1R7
                                                              linear
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                                            STRANDEDNESS:
TOPOLOGY: line
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INFORMATION FOR
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                                                                                        US-09-293-533-16
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CITY: T
         SEQUENCE CI
LENGTH:
TYPE: AT
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; Sequence 4, Application US/08973518
; Patent No. 6328962
; Patent No. 6328962
; Patent No. 6328962
; APPLICANT: Anand, Naveen N
; APPLICANT: Cates, George A
; APPLICANT: Caterini, Judith E
; APPLICANT: Caterini, Judith E
; APPLICANT: Caterini, Michel H
; TITLE OF INVENTION: CHIMERIC ANTIBODIES FOR DELIVERY OF
; TITLE OF INVENTION: ANTIGENS TO SELECTED CELLS OF THE IMM
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Sim & McBurney
; ADDRESSEE: Sim & McBurney
                                                             QVQLQESGPGLVAPSQSLSITCTVSGISLNRYGVHWVRQPPGKGLEWL
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; DB 3;
..8e-47;
les 13;
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 Score 531.5;
Pred. No. 1.8e
4; Mismatches
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,518
FILING DATE: 07-APR-1998
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1038-781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                      4, Application US/08973518
>. 6328962
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ilarity 85.1%;
Conservative
 83.7%;
ilarity 85.1%;
Conservative
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTIC LENGTH: 140 amino ac
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Best Local Similarity
Matches 103; Conser
                 Local Similarity
nes 103; Conser
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Ontario
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PRIOR APPLICATION DATA:
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                                                                                                                                                ; TOPOLOGY:
US-08-881-037-67
                                               TELEFAX:
TELEX:
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Best Local S
Matches 104
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                                                                                                                     KARL
                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                              ;Patent No. 5455030
; APPLICANT: LADNER, ROBERT C.;BIRD, ROBERT E.;HARDMAN,
; TITLE OF INVENTION: IMMUNOTHERAPHY USING SINGLE CHAIN
;POLYPEPTIDE BINDING MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                     Score 529.5; DB 6;
Pred. No. 5.7e-47;
5; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-881-037-67

Sequence 67, Application US/08881037

Patent No. 6080588

GENERAL INFORMATION:

APPLICANT: Swanson, Patrick C.

TITLE OF INVENTION:

CORRESPONDENCES: 113

CORRESPONDENCES: 113

ADDRESSEE: Morrison & Foerster

STREET: 755 Page Mill Road

CITY: Palo Alto

STATE: CA

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version
                                                                                                                                                   NUMBER OF SEQUENCES: 24

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/40,440
FILING DATE: 1-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 512,910
FILING DATE: 25-APR-1990
APPLICATION NUMBER: 299,617
FILING DATE: 19-JAN-1989
APPLICATION NUMBER: 92,110
FILING DATE: 02-SEP-1987
APPLICATION NUMBER: 902,971
FILING DATE: 01-SEP-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MBER: US 08/443,540
18-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/881,037
FILING DATE: 23-JUN-1997
CLASSIFICATION: 530
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larity 85.8%;
Conservative
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 18-MAY-
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103; Conserv
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LENGTH:
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Best Local S
Matches 103
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121
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Mediated Disorder
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APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Chok, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Ustrite OF INVENTION: Treatment of IL5 Mediated Dison NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/667,769A
                                                                                                                                                                                                                                                                                              14;
                                                                                                                                                                                                                                                              DB 3;
                                                                                                                                                                                                                                                                             .4e-47;
                                                                                                                                                                                                                                                            Score 525.5;
Pred. No. 6.4e
2; Mismatches
REGISTRATION NUMBER: 34,202
REFERENCE/DOCKET NUMBER: 203442110710
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEFAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MBER: PCT/US95/17082
22-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MBER: US 08/467420
06-JUN-1995
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                                                                                                                                                                                                                                                                                            2 ;
                                                                                                              67
                                                                                                                                                                                                                                                          ch 82.8%;
il Similarity 86.0%;
104; Conservative
                                                                                                                                                121 amino acids amino acids
                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 121 amino acid
TYPE: amino acid
                                                                                                                                                                             STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
FILING DATE: 22-DEC-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15, Applicat
Patent No. 5783184
GENERAL INFORMATION:
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APPLICATION NUMBER:
FILING DATE: 23-DE
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ZIP: 02142
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TYPE: a
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د.
                                                                                                                                                                                                                                                                 Length 119
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APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant ILS Antagonists Useful i
                                                                                                                                                                                                                                                                                          Indels
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ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                               16;
12;
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STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                              Score 523; DB 1;
Pred. No. 1.1e-46
7; Mismatches 12
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       US 08/363131
                                                                             50503
              FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P505
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 15:
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FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                 82.4%;
                                                                                                                                                               119 amino acids
amino acid
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                                                                                                                                                                                     STRANDEDNESS: single
TOPOLOGY: linear
DLECULE TYPE: protein
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CORRESPONDENCE ADDRESS
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        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ESULT 12
CT-US95-17082A-15
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US-08-667-769A-15
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APPLICANT: Benjamin, Christopher D
APPLICANT: Hession, Catherine A
APPLICANT: Whitty, Adrian
TITLE OF INVENTION: COMMON GAMMA CHAIN BLOCKING AGENTS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Blogen 1--
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Pred. No. 1.1e-46
7; Mismatches 1
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                                                  NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: A006 PCT CIP
FILING DATE: 09-MAY-1997

APPLICATION NUMBER: 60/017,466

FILING DATE: 10-MAY-1996

ATTORNEY/AGENT INFORMATION:

NAME: Kaplan, Warren A.

REGISTRATION NUMBER: 34,199

REFERENCE/DOCKET NUMBER: A006 PCT (TELECOMMUNICATION INFORMATION:
TELEPHONE: 617 679-2000

TELEFAX: 617 679-2838
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US/09/189,129
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
US 08/363131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3, Application US/09189129
Patent No. 6323027
GENERAL INFORMATION:
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Cambridge Center
                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 82.6%;
Matches 100; Conservative
               FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                       119 amino acids
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OPERATING SYSTEM:
SOFTWARE: Patent
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CITY: Cambridge
STATE: Massachus
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TOPOLOGY: line
MOLECULE TYPE:
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Best Local Similarity
Matches 100; Conser
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US-08-860-174A-2
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US-08-860-174A-2
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/824,286
FILING DATE: 02-Apr-2001
CLASSIFICATION - CUNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/017,466
FILING DATE: 10-MAY-1996
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A006 PCT CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617 679-2000
TELEFAX: 617 679-2838
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                             Score 521.5; DB 3;
Pred. No. 1.5e-46;
3; Mismatches 6;
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Pred. No. 1.5e-46;
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Burkly, Linda C
Benjamin, Christopher D
Hession, Catherine A
Whitty, Adrian
TITLE OF INVENTION: COMMON GAMMA CH
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Biogen, Inc.
STREET: 14 Cambridge
CITY: Cambridge
STATE: Massachusetts
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/ FRAGMENT TYPE: internal
/ SEQUENCE DESCRIPTION: SEQ ID NO: 3
US-09-824-286-3
                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/09824286
Patent No. 6770745
GENERAL INFORMATION:
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M
                                                                                                                            th 82.1%;
Similarity 85.2%;
98; Conservative
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internal
NO:
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 V FOR SEQ ID NO:
CHARACTERISTICS
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                                                               linear
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FRAGMENT TYPE:
                                                 STRANDEDNESS
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LENGTH:
TYPE: at
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INFORMATION
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Sequence 2, Application US/08860174A;
Patent No. 5989830;
GENERAL INFORMATION:
APPLICANT: DAVIS, Paul James
APPLICANT: VERHOEIJEN, Martine Elisa
APPLICANT: WILSON, Steve
TITLE OF INVENTION: A BIFUNCTIONAL OR BIVALENT
TITLE OF INVENTION: ANTIBODY FRAGMENT ANALOGUE
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
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N.W.
Mismatches
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS DOS
SOFTWARE: MS WORD, WINDOWS 95, Ver. 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,174A
FILING DATE: June 16, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER: EP 95307332.7
October 16, 1995
NUMBER: PCT/EP/96/03605
August 14, 1996
SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2004, 13:01:10
                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: PILLSBURY, MADISON
STREET: 1100 New York Avenue,
STREET: 9th Floor, East Tower
CITY: WASHINGTON, D.C.
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82.6%;
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amino acid
Y: linear
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APPLICATION NUMBER: EP
FILING DATE: October 16
APPLICATION NUMBER: PCT
FILING DATE: August 14,
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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US-09-423-439-58
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US-08-190-199A-65
US-08-956-047-33
US-08-423-439-44
US-08-467-393-4
US-08-483-528B-92
US-08-438-562-2
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US-08-434-122-33
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  OTHER INFORMATION:
9-171-945-17
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Percent Similarity:
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Matches:
Conservative:
Mismatches:
Indels:
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ZIP: 19103

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                             Antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0032
TELECOMMUNICATION INFORMATION:
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                          US/08/303,569B
                                                                                                                  sequence 5, Application US/08303569B; Patent No. 5859205; GENERAL INFORMATION:
APPLICANT: Adair, John R.
APPLICANT: Athwal, Diljeet S.
APPLICANT: Emtage, John S.
TITLE OF INVENTION: Humanised Ant NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kustreet: One Liberty Place - 46t.
                                                                                                                                             US/08303569B
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                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 07-SEP-19
CLASSIFICATION:
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Best Local Similarity:
Query Match:
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5455030-3
; Patent No. 5455030
; TITLE OF INVENTION: IMMUNOTHERAPHY US; POLYPEPTIDE BINDING MOLECULES
; NUMBER OF SEQUENCES: 24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/40,440
; TITLE OF SEQUENCES: 24
                  -09-171-945
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                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 512,910
FILING DATE: 25-APR-1990
APPLICATION NUMBER: 299,617
FILING DATE: 19-JAN-1989
APPLICATION NUMBER: 92,110
FILING DATE: 02-SEP-1987
APPLICATION NUMBER: 902,971
FILING DATE: 01-SEP-1986
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US-09-795-515-5
; Sequence 5, Application US/09795515
; Patent No. 6632927
; GENERAL INFORMATION:
; APPLICANT: Athwal, Diljeet S.
APPLICANT: Emtage, John S.
; TITLE OF INVENTION: Humanised Ant;
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Woodcock Washburn Ku
; STREET: One Liberty Place - 46t
; STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
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Best Local Similarity
Query Match:
DB:
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US-09-795-515-5
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 AGTGGCAGTGGGTCTGGGACAGAATTCACCCTGGAAATCAGTAGAGTGAA
                                                                                                                                                                             Sequence 5, Application US/08116247;
Patent No. 5929212;
GENERAL INFORMATION:
APPLICANT: Jolliffe, Linda K.
APPLICANT: Adair, John R.
APPLICANT: Athwal, Diljeet S.
TITLE OF INVENTION: CD3 Specific Recombinant Antibody
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Wasburn Kurtz Mackiewicz & No. 5929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 235
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    One Liberty Place - 46th Floor
Philadelphia
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Matches:
Conservative:
Mismatches:
Indels:
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R: CARP-0011
                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/116,24'FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07/743,37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 235 amino acids
TYPE: amino acids
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 10-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Paintin, Francis A.
REGISTRATION NUMBER: 19,38
REFERENCE/DOCKET NUMBER: C
                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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75.00%
66.02%
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US-08-116-247-5
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CITY: P
STATE:
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Conservative:
Mismatches:
Indels:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vere
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,515
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION
OPERATING DATE:
APPLICATION NUMBER: 08/846,658
FILING DATE: 01-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0057
TELECOMMUNICATION INFORMATION:
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RESULT
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; Sequence 5, Application US/09348224
; Patent No. 6750325
; GENERAL INFORMATION:
; APPLICANT: Jollieffe, Linda Kay
APPLICANT: Adair, John Robert
; APPLICANT: Athwal, Diljeet Singh
; TITLE OF INVENTION: CD3 Specific Recominant
; FILE REFERENCE: CARPO066
; CURRENT APPLICATION NUMBER: US/09/348,224
; CURRENT FILING DATE: 1999-07-06
; EARLIER APPLICATION NUMBER: 08/116,247
; EARLIER FILING DATE: 1991-10-04
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: PatentIn Ver. 2.1
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Matches:
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Query Match:
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US-09-553-498-8
; Sequence 8, Application US/09553498
; Patent No. 6309861
; GENERAL INFORMATION:
; APPLICANT: Ambrosius, Dorothee
; APPLICANT: Rudolph, Rainer
; APPLICANT: Schwarz, Elisabeth
; TITLE OF INVENTION: Process for the production of nat
; FILE REFERENCE: Case 20379
; CURRENT APPLICATION NUMBER: US/09/553,498
; CURRENT FILING DATE: 2000-04-20
; PRIOR APPLICATION NUMBER: EP99107412.1
; PRIOR FILING DATE: 1999-04-26
; NUMBER OF SEQ ID NOS: 10
; SEQ ID NO 8
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Matches:
Conservative:
Mismatches:
Indels:
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; Sequence 8, Application US/09618869; Patent No. 6455279
; GENERAL INFORMATION:
; APPLICANT: Ambrosius, Dorthee
; APPLICANT: Rudolph, Rainer
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Rudolph, Rainer
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; TYPE: PRT
; ORGANISM: E. coli
US-09-553-498-8
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Best Local Similarity
Query Match:
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Batent No. 5885816

GENERAL INFORMATION:

TITLE OF INVENTION:

TITLE OF INVENTION:

TITLE OF INVENTION:

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSE: Wenderoth, Lind & Ponack

STREET: 805 Fifteenth Street, N.W., #700
APPLICANT: Schaeffner, Joerg
APPLICANT: Schwarz, Elisabeth
TITLE OF INVENTION: PROCESS FOR THE PRODUCTION OF NATURALLY I
TITLE OF INVENTION: CHAPERONES
FILE REFERENCE: 20381
CURRENT APPLICATION NUMBER: US/09/618,869
CURRENT FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: EP99114811.5
PRIOR FILING DATE: 1999-07-29
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Matches:
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Mismatches:
Indels:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 i.
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordperfect 5.1
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444.00
86.36%
80.00%
65.88%
                                                                                                                                                ORGANISM: Escherichia col-09-618-869-8
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                                                                                                 NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver.
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U.S.A.
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Best Local Similarity
Query Match:
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TYPE: PRT
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Sequence 9, Application US/09171945
Sequence 9, Application US/09171945
Batent No. 6277599
GENERAL INFORMATION:
APPLICANT: Emery, Stephen
APPLICANT: Copley, Clive Graham
APPLICANT: Edge, Michael Derek
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said
TITLE OF INVENTION: Antibody, and Their Therapeutic Use in an Adept System
FILE REFERENCE: Monoclonal Antibody to CEA
CURRENT APPLICATION NUMBER: US/09/171,945
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: GB9703103.3
PRIOR FILING DATE: 1997-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACCTICCCCAAAAGAAGGATITAIGACACAICCAAACIGGCIICIGGAGICCCIGCICGC
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Matches:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/737,129A
FILING DATE: No. 5885816ember 15, 1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels:
Gaps:
                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 213 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                single
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Best Local Similarity:
Query Match:
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Matches:
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Mismatches:
Indels:
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                                                                                                                                                                                                                                      RESULT 12
US-09-526-738A-4
; Sequence 4, Application US/09526738A
; Patent No. 6630584
; GENERAL INFORMATION:
; APPLICANT: LTD.
; TITLE OF INVENTION: SINGLE CHAIN ANTIBODY AG
; FILE REFERENCE: 1196336
; CURRENT APPLICATION NUMBER: US/09/526,738A
; CURRENT FILING DATE: 2000-03-16
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 4
; LENGTH: 258
; TYPE: PRT
; ORGANISM: Humanus
; ORGANISM: Humanus
US-09-526-738A-4
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/09526738A

Patent No. 6630584

GENERAL INFORMATION:
APPLICANT: RAMOT UNIVERSITY AUTHORITY FOR APPLIED RESEARCH
APPLICANT: LTD.
TITLE OF INVENTION: SINGLE CHAIN ANTIBODY AGAINST MUTANT F
FILE REFERENCE: 1196336
CURRENT APPLICATION NUMBER: US/09/526,738A
CURRENT FILING DATE: 2000-03-16
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
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Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
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                                                                                             Artificial
                                                                                                                               Length:
Matches:
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                                                                                                                                                                        Gaps:
PRIOR APPLICATION NUMBER: GB9609405.7
PRIOR FILING DATE: 1996-05-04
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR FILING DATE: 1997-04-29
NUMBER OF SEQ ID NOS: 131
SOFTWARE: Patentin Ver. 2.1
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                                                                                               ; OTHER INFORMATION: Description US-09-171-945-9
                                                                              Sequence
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439.00
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427.00
80.70%
73.68%
63.35%
                                                   SEQ ID NO 9
LENGTH: 108
TYPE: PRT
ORGANISM: Artificial S
FEATURE:
                                                                                                                                                                                          US-10-049-868A-1 (1-359)
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Best Local Similarity:
Query Match:
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Best Local Similarity
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.09-526-738A-2
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Query Match:
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                                                                                                                                             APPLICANT: Yashwant M. Deo
APPLICANT: Joel Goldstein
APPLICANT: Robert Graziano
APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHTUF
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,052
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Mismatches:
Indels:
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Matches:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/484,172
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI-043CP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                          SSEE: LAHIVE & COCKFIELD
T: 60 State Street, Suite
Boston
                                                                                                                             US-08-661-052-16; Sequence 16, Application US/08661052; Patent No. 5837243; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
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426.00
85.71%
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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US-08-661-052-16
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CITY: BO:
STATE: Ma
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GATGTGGGTGTATTACTGTCAACAACTTGTAGAGTATCCGCTCACGTT
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                                             8th Floor
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Mismatches:
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                                                                                                                     Sequence 2, Application US/08652507
; Patent No. 5876691
; GENERAL INFORMATION:
   TITLE OF INVENTION:
   TITLE OF INVENTION:
   NUMBER OF SEQUENCES: 8
   CORRESPONDENCE ADDRESS:
   ADDRESSEE: Nixon & Vanderhye, P.C.
   STREET: 1100 No. 5876691th Glebe Road, 8t
   CITY: Arlington
   STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels:
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ZIP: 22201-4,1
ZIP: 22201-4,1
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,507
FILING DATE: 02-Jul-1996
FILING DATE: 530
                                                                                                                                                                                                                                                                                                                                                                                    AFFLICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Arthur R. Crawford
REGISTRATION NUMBER: 25,327
REFERENCE/DOCKET NUMBER: 117-211
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4100
TELEFAX: 703-816-4100
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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85.71%
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ZIP: 22201-4714
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Best Local Similarity:
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US-08-652-507-2
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US-09-188-082-16
; Sequence 16, Application US/09188082
; Patent No. 6270765
; Patent No. 6270765
; Patent No. 6270765
; APPLICANT: Yashwant M. Deo
; APPLICANT: Acel Goldstein
; APPLICANT: Chezian Somasundaram
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS COMPRISED
TITLE OF INVENTION: OF ANTI-FC RECEPTOR ANTIBODIES
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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Matches:
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APPLICATION NUMBER: 08/661,052
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Arnold, Beth E.
REGISTRATION NUMBER: 35,430
REFERENCE/DOCKET NUMBER: MXI~043CP
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60 State Street, Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 553 amino acids
TYPE: amino acids
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OPERATING SYSTEM: PC-DOS
SOFTWARE: Patentin Relea
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/O
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MOLECULE TY
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GluLeuLysArgAla 5
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